Hart, Edward

From:

Subject:

Swope, Sheridan

Sent:

Friday, March 05, 2004 11:16 AM

To:

Hart, Edward FW: 09/966,880

Importance:

High

Ed, the applicants have again made changes to the claims that require an additional search for this case (we're negotiating an allowance).

So, would you add the following to my rush search? For 09/966,880, Pls search and interference search:

SID 7: full-length and oligo search (20NTs) against the NT data bases.

Thanks in advance for your great service! Sheridan

[Swope, Sheridan] Sheridan Swope, Ph.D. Patent Examiner, AU 1652
Recombinant Enzymes
571-272-0943 (voice)
E03A70 Remsen Bld (Office)
E03C70 Remsen Bld (Mailbox)

1951 /vol



STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 115892

TO: Sheridan Swope Location: REM-3A70

Art Unit: 1652

Friday, March 05, 2004

Case Serial Number: 09/966880

From: Edward Hart

Location: Biotech-Chem Library

REM-1A55

Phone: 571-272-2512

edward.hart@uspto.gov

Search Notes

Examiner Swope,

Here are the results of the search you requested.

Please feel free to contact me if you have any questions.

Edward Hart



4.4	Database :	ocessing	um DB seq	Word size : Total number of h	Searched: 3	Scoring table: 0	Title: U Perfect score: 5 Sequence: 1	Run on:	OM nucleic - nucleic	
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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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ALIGNMENTS

AUTHORS TITLE	REFERENCE	SOURCE ORGANISM	KEYWORDS	ACCESSION	DEFINITION	Locus	RESULT 1
Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A. Method and reagent for treating diseases or conditions concerning	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 17)	Homo sapiens (human) Homo sapiens	JP 2002509721-A/3934.	BD200908 T. 33010678	Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response.	BD200908 17 bp RNA linear PAT 17-JUL-2003	

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OS HOMO Sapiens (human)
PN JP 2002509721-A/3934
PD 02-APR-2002
PF 24-MAR-1999 JP 2000541291
PR 27-MAR-1998 US 60/079678
PI PAMELA A PAVCO, ELISABETH ROBERTS, THALE JARVIS
PI JAMES A MCSWIGGEN
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/mol_type="genomic RNA"
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Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response.

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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequences involved in tumoral suppression, a
apoptosis and/or viral resistance phenomena
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Catarrhini; Hominidae;
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7e+04;
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. 7e+04;
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resistance and their use
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Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

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OS Homo sapiens (human)
PN JP 2002509721-A/3935
PD 02-APR-2002
PF 24-MAR-1999 JP 2000541291
PF 27-MAR-1998 US 60/0796
PI PAMELA A PAVCO, ELISABETH I
PI JAMES A MCSWIGGEN
PC
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PC A61P35
C12N5/00
                                                                                                                                                                                    Miraglia,L., Bennett,C.Frank., Dean,N. Antisense compositions and methods for interleukin-1 receptor expression Patent: US 5856099-A 1 05-UAN-1999;
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AR026494
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27-MAR-1998 US 60/079678
PAMELA A PAVCO,ELISABETH ROBERTS,THALE JARVIS,CLAIRE COESHOTT,
                                                         Conservative
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                                                                                                                          /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="genomic RNA"
/db_xref="taxon:9606"
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100.0%; Pred. No.
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RESULT 7 AR026495

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Hordeum vulgare
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Pooideae; Triticeae; Hordeum.
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Miraglia,L., Bennett,C.Frank., Dean,N.
Antisense compositions and methods for
interleukin-1 receptor expression
Patent: US 5856099-A 2 05-UAN-1999;
                                                                                                     AX323965
Sequence 103 from Patent
AX323965
                                                                                                                                                                                                                                                                                                                                                                                 Patent: WO 02053728-A 4 Elitra Pharmaceuticals,
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Candida albicans
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                                                                                                                                                                                                                                                                                                                  /organism="Candida albicans"
/mol_type="unassigned DNA"
/db_xref="taxon:5476"
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AX323966
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hordeum vulgare
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Pooideae; Triticeae; Hordeum.
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Targeted chromosomal genomic alterations in plants using modified single stranded oligonucleotides
Patent: WO 0192512-A 103 06-DEC-2001;
UNIVERSITY OF DELAWARE (US)
                                                                                                                                                  Sequence 795 from Patent AX499488 AX499488.1 GI:23381781
                                   Human testis expressed patched like protein Patent: EP 1229046-A 795 07-AUG-2002;
                                                                                                             Homo sapiens (human)
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                                                               Zhan, J.
                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
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Location/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:4513"
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/mol_type="unassigned DNA"
/db_xref="taxon:4513"
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.00.0%; Pred. No. 2.5e+05;
.ve 0; Mismatches 0;
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Primates;
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                                                                                      Craniata; Vertebrata;
Catarrhini; Hominidae;
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385 CTGCACCGCGCCG 397
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13; Conserv
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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AX499492/c
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1 Similarity 100.0%; Pred. No. 2.5e+05;
13; Conservative 0; Mismatches 0;
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Sequence 799 from Patent
AX499492
AX499492.1 GI:23381785
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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        17 bp DNA
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Sequence 314
AX687582
AX687582.1
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Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumour suppression, tumour suppression, apoptosis and/or resistance to viruses and their
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                           CICCIGGAGCCCC 5
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314 from Patent EP1281758
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                              GI:29410278
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thes 0;
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mdz4, mdz7

Euteleostomi;

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RESULT 20
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AX687583/c
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Sequence 316 from Patent
AX687584
AX687584.1 GI:29410280
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Sequence 315 from Patent
AX687583
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Patent: EP 1281758-A 316 05-FEB-2003;
Aeomica, Inc. (US)
                                           Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins
                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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AX735945/c
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AX687585/c
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                                                                                          thereof as medicaments
Patent: WO 03025177-A 1535 27-MAR-2003;
Molecular Engines Laboratories (FR)
                                                                                                                                    Telerman, A., Amson, R. and Tuijnder, M. Sequences involved in phenomena of tumour reversion, apoptosis and/or resistance to
                                                                                                                                                                                                 Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Similarity 100.0%; Pred. No. 2.5e+05;
13; Conservative 0; Mismatches 0;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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100.0%; Pred. No. 2.5e+05;
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suppression, tumour viruses and the use

PAT 08-MAY-2003

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Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: US 6468983-A 9 22-OCT-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 bp

Kit and method for determining

BD104902

BD104902.1 GI:22650476

WO 0132572-A/1006.
                                                                                                                                                                                                                                                                                                     Sequence 9
AR241182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kit and method for determining HLA type
Patent: WO 0192572-A 1006 06-DEC-2001;
NISSHINBO INDUSTRIES INC,SYSTEM RESEARCH INC,HIDETOSHI INOKO, TR
KAGIYA, TATSUO ICHIHARA,YOSHIYUKI MATSUMURA,SHOGO MORIYA,MICHIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         artificial sequences.

1 (bases 1 to 17)
1 (bases 1, to 18)
1 (bases 1
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synthetic construct
                                                                                                                                                                                                                       Unknown.
                                                                                                                                                                                                                                                                             AR241182.1
                                                                                                                                                                      Unclassified.
                                                                                                                                                                                                 Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SHOGO MORIYA, MICHIO NISHIDA
C12Q1/68,C12M1/00,C12N15/09,G01N33/53
Description of Artificial Sequence:capture
Key Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-2001 WO 2001JP004662
01-JUN-2000 JP 00P 164798
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WO 0192572-A/1006
06-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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100.0%; Pred. No.
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6468983.
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J. 2.5e+05;
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                                                                                                                                                                                                                                                                                                                                                     DNA
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AUTHORS
TITLE
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AX129413/c
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KEYWORDS
SOURCE
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AR253259/c
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Matches 13
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TITLE
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                        Homo sapiens (human)
                                                                                                                                                                                                                                                                        Sequence 631 from Patent AX129413
                                                                                                                                                      diseases
                                                                                                                                                             Robbins, J.M. and Tritz, R.
Ribozyme therapy for the treatment of
                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                           AX129413.1 GI:14135718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Non-stochastic generation of genetic vaccines
Patent: US 6479258-A 4 12-NOV-2002;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unknown
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                                                                                                                                        Patent: WO 0130362-A 631
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                                                                   /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mol_type="genomic
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/mol_type="genomic DNA"
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Score 13; DB Pred. No. 2.5 O; Mismatches
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Pred. No.
                                                                                                                                        03-MAY-2001;
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6479258.
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                                                        binding site"
          DB 6; L
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2.5e+05;
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                    Length 19;
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AX259857/c
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AX129414/c
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Best Local (
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AX129414
                                                                                                                                                                                                          Hugot, J.P., Thomas, G., Zouali, M., Lesage, S
Genes involved in intestinal inflammatory
Patent: WO 0172822-A 84 04-0CT-2001;
Fondation Jean Dausset-Ceph (FR)
                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Robbins, {\tt J.M.} and {\tt Tritz,R.} Ribozyme therapy for the treatment of
                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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 BD084645
                                                                                                                                                                                                                                                                                                 Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent: WO 0130362-A 632 03-MAY-2001;
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Mammalia; Eutheria; Primates;
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                                                                                                                                                  /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note="Cdk6 ribozyme binding"
                                                                                                                                                                                                ocation/Qualifiers
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J. 2.5e+05;
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                                                                                                                   Length 19;
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PAT
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27-AUG-2002
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PR 21-APR-1997 US 60/044507,03-FEB-1998 ROBERT H SILVERMAN,SEIJI KONDO,JOHN K COWELL,(PI TORRENCE PC C07H21/00,C07H21/02.C1??
Seattle, WA 98104, ....
Seattle, WA 98104, ....
e-mail: EAOstrander@lbl.gov
PCR Buffer: PCR buffer (Perkin-Elmer/Cetus)
PCR Profile: Denaturation: 94 degrees C for 1.00 minute
PCR Profile: Denaturation: 94 degrees C for 0.45 minutes
Annealing: 74 degrees C for 1.00 minutes
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13; Conserv
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                                                                                                                                                                                                                                                                       Transplantation Biology Dept
1124 Columbia; Mailstop M318
Seattle, WA 98104, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                        Original source text: Canis familiaris (library: pBluescript+) adult spleen DNA.
Submitted by:
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0 (strander,E.A., Mapa,F.A., Yee,M. and Rine,J.
0 (one hundred and one new simple sequence repeat-based markers for the canine genome (strander) (s
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Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
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1 (bases 1 to 19)
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Description of Artificial Sequence: oligonucleotide
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/db_xref="taxon:32630"
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179708
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Trofatter, J.A., MacCollin, M.M. and Gusella, J.F.
Tumor suppressor merlin and annibodies thereof
Patent: US 6077685-A 3 20-UN-2000;
Location/Qualifiers
                                                                           1 (bases 1 to 20)
Trofatter, J.A., MacCollin, M.M. and Gusella,
Tumor suppressor gene merlin
Patent: US 5707863-A 3 13-JAN-1998;
Location/Qualifiers
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Final Extension:
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/moi_type="unassigned DNA"
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/dev_stage="adult"
/tissue_lib="B. Ostrander,
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sinsion: 74 degrees
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/mol_type="genomic DNA"
/db_xref="taxon:9615"
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AR257223
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Patent: US 6566135-A 100 20-MAY-2003;
Location/Qualifiers
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/mol_type="genomic
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/mol_type="genomic DNA"
                                   from Patent W00233056.
                  GI:21538245
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100.0%; Pr
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Pred. No.
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2.5e+05;
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2.5e+05;
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Homo sapiens (human)

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SOURCE
ORGANISM
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VERSION
KEYWORDS
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BD138115/c
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DEFINITION
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1 (bases 1 to 20)

TITLE
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 41 26-MAR-2002;

MENT
OS
Unidentified
PD 26-MAR-2002
PF 26-MAR-2002
PF 26-MAR-2002
PF 26-MAR-2002
PF 26-MAR-2009
                                                                                                                                                                                                                                                                                                                           COMMENT
                                                                                   ORIGIN
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TITLE
JOURNAL
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TITLE
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Best Local Similarity
Matches 13; Conserv
                                                   Query Match
Best Local :
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Antisense modulation of human
BD138115
BD138115.1 GI:23233060
JP 2002508944-A/41.
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Eukaryota; Metazoa;
                                                                                                                                                                               Mammalia; Eutheria;
                                                                                                                                                  44
                                                    Similarity
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 GAGCCCGAGGGGC 7
                   GAGCCCGAGGGGC 376
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ilarity 100.0%;
Conservative
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C12N15/00
                                       2.2%;
ilarity 100.0%;
Conservative
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Topology: Linear;
Antisense modulation of human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="PCR primer"
                                                                                           /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                     /organism='Unidentified'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="unassigned
/db_xref="taxon:9606"
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3. 2.5e+05;
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KEYWORDS
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AX358112/c
LOCUS
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AUTHORS
TITLE
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I43304/c
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RESULT 39
A09427/c
LOCUS
DEFINITION
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Oligonucleotide (b3).
A09427
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Arabidopsis thaliana
Arabidopsis thaliana
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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AX358112
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Szostak, J. W. and Huizenga, D.E.
DNA aptamers and catalysts that bind adenosine or adenosine-5'-phosphates and methods for isolation Patent: US 5631146-A 122 20-MAY-1997;
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143304
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synthetic construct
synthetic construct
artificial sequences
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                                                                                                                                                                                                  CTTCCTCCGCTA 191
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nilarity 100.0%;
Conservative 0
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/mol_type="unassigned DNA"
/db_xref="taxon:3702"
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                                                                                                                                                                                                                                                                                                       /organism="unknown"
/mol_type="unassign
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from Patent W00194394.
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A10630/c
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                                                                                                     synthetic construct synthetic construct artificial sequences.

1 (bases 1 to 15)

1 (bases 1 to 15)

1 Ueda, I., Niwa, M., Saito, Y., Sato, S., Ono, H. and Kitaguchi, T.

59 Valine insulin-like growth factor I and process for production
                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
12; Conserv
                                                                                                                                                                                                                             oligonucleotide 'b3'
A11578
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Oligonucleotide (B3).
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                                                                      Patent: EP 0158892-A 74 23-OCT-1985; FUJISAWA PHARMACEUTICAL CO., LTD
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                            l Similarity
12; Conserv
                                                                                                                                                Ueda,I., Niwa,M., Satoh,S., Saitoh,Y. Process for production of isulin-like for production thereof Patent: EP 0219814-A 48 29-APR-1987; FUJISAWA PHARMACEUTICAL CO., LTD Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                           3 CTCCTGGAGCCC 14
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Location/Qualifiers
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1 (bases 1 to 15)
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llarity 100.0%; Pred. No. 9.1e+05;
Conservative 0; Mismatches 0;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                   organism="synthetic construct"
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/db_xref="taxon:32630"
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Sequence 9 from Patent WO9801757.
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Sequence 33 from Patent WO9744355.
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Windass, J.D., Duncan, R.E., Baule, V.J.
TOXINS FROM THE WASP BRACON HEBETOR
Patent: WO 9616171-A 8 30-MAY-1996;
ZENECA LTD (GB)
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unidentified unidentified
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Duncan,R.E., Suner,M., Daly,A., Christian,P.D.,
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type≈"unassigned DNA"
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/mol_type="unassigned DNA"
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Patent: US 5747645-A 10 05-MAY-1998;
Location/Qualifiers
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Osbourn,J.K., Derbyshire,E.J., Mccafferty,J.G.,
Johnson,K.S.
LABELLING AND SELECTION OF MOLECULES
Patent: WO 9801757-A 9 15-JAN-1998;
CAMBRIDGE ANTIBODY TECH (GB)
Other publication GB 2315125 19980121.
1 (bases 1 to 15)
Osbourn, J. Katharine., Derbyshire, E. Joy.,
Vaughan, T. John. and Johnson, K. Stuart.
Labelling and selection of molecules
Patent: US 5994519-A 35 30-NOV-1999;
Location/Qualifiers
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I28574
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AR126265
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Watanabe, K.A., Ren, W.-Y. and Weil, R.
Complementary DNA and toxins
Patent: US 5571937-A 27 05-NOV-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unknown
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 I58736
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/mol_type="unassigned DNA"
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I58736
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Sprecher, C.A. cod
Cytoplasmic antiproteinase-2 and cod
Patent: US 5712117-A 10 27-JAN-1998;
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Patent: US 5710026-A 10 20-JAN-1998;
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Sequence 10 from patent US 5710026.
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Watanabe, K.A., Ren, W.-Y. and Weil, R.
Complementary DNA and toxins
Patent: US 5652350-A 27 29-JUL-1997;
                                                                         Unclassified.
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/mol_type="unassigned
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Vaughan, T. John. and Johnson, K. Stuart.
Labelling and selection of molecules
Patent: US 6342588-A 35 29-JAN-2002;
                                                                                         Similarity
12; Conserv
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Osbourn,J.K., Derbyshire,E.J., McCafferty,J.G.,
Johnson,K.S.
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/mol_type="unassigned
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from patent US 6342588.
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INTERNATI REAGENTS CORP

OS None

OC Artificial sequences.

PN JP 1994133799-A/4

PN 19-1994133799-A/4

PF 27-OCT-1992 JP 1992311416

PI YAMANISHI KOICHI, YAMAMOTO TAKESHI, MORI HIROYUKI PC C12Q1/68,C12Q1/68,C12N15/11,C12N15/38;

CC strandedness: Single;
CC typology: Linear;
CC hypothetical: No;
CC anti-sense: No;
FH Key Location/Qualifiers
FH source 1..16
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FT source 1...15
FT fource /organism='Artificial Sequence'.
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Patent: WO 0159107-A 18 16-AUG-2001;
FUSO PHARMACEUTICAL INDUSTRIES LTD,NOBUTAKA WAKAMIYA
                                                                                                                                                                                             1 (bases 1 to 16)
Yamamishi,K., Yamamoto,T. and Mori,H.
Yamamishi,K., Yamamoto,T. and Mori,H.
ANALYSIS OF HUMAN HERPES VIRUS 6 TYPE @(3754/24)HHV-6) DNA AND
DISCRIMINATION OF SUB-TYPE
Patent: JP 1994133799-A 4 17-MAY-1994;
                                                                                                                                                                                                                                                                             unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-FEB-2001 WO 2001JP000874
14-FEB-2000 JP 00P 35155,10-OCT-2000 JP
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Location/Qualifiers
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PAT 29-SEP-1997

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AR361170
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                                                                                                                      Stinchcomb, D.T. and McSwiggen, J.A.
Interleukin-2 receptor gamma-chain ribozymes
Patent: US 5807743-A 1283 15-SEP-1998;
Location/Qualifiers
                                                                                                                                                                                                                          Sequence 1283 from patent AR040435
                                                                                                                                                                                                                                                                                                                                                                                                                                 Methods for detection of transition single-nucleotide polymorphisms Patent: US 6599700-A 41 29-JUL-2003;
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Bellacosa, A.
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                                                                                       /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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490 GGGCTGCATGAA 501
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Sidransky, D.
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                                                                              /mol_type="unassigned
                                                                                                                                                                                                       GI:1606706
                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
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/db_xref="taxon:9606"
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

CE 1 (bases 1 to 17)

RS Landers, J.E., Jordan, B., Housman, D.E. and Charest, A.

Methods and products related to genotyping and DNA analysis

Patent; JP 2002525127-A 367 13-AUG-2002;

MASSACHUSETTS INSTITUTE OF TECHNOLOGY

OS Homo sapiens (human)

PN JP 2002525127-A/367

PD 13-AUG-2002

PP 13-AUG-2002

PP 24-SEP-1998 US 60/101757

PR 25-SEP-1998 US 60/101757

PI JOHN E LANDERS, BARBARA JORDAN, DAVID E HOUSMAN, ALAIN CHAREST PC C12N15/09, C12Q1/68, G01N33/53, G01N33/566, G01N33/58, G01N37/00, PC G01N37/00, C12Q1/68, G01N33/53, G01N33/566, G01N33/58, G01N37/00, PC CC Methods and products related to genotyping and DNA analysis FH
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Y Location/Qualifiers
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Sequence 159 from patent US 5567809.
127987
Unclassified.
1 (bases 1 to 17)
Sidransky,D.
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Apple,R.J., Erlich,H.A., Griffith,R.L. and Scha
Methods and reagents for HLA DEDeta DNA typing
Patent: US 5567809-A 138 22-OCT-1996;
                                                                               Sequence 59 from patent US 191577
                                                                                                                                                                                                                                                                                                                1 (bases 1 to 17)
Apple, R.J., Erlich, H.A., Griffith, R.L. and Sch.
Methods and reagents for HLA DRbeta DNA typing
Patent: US 5567809-A 159 22-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                       Unknown
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/mol_type="unassigned
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/mol_type="unassigned DNA"
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Pred. No. 9.1e+05;
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AX214609/c
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AX215505/c
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Patent: US 5726019-A 59 10-MAR-1998;
Location/Qualifiers
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                                          nogo gene expression
Patent: WO 0159103-A 947 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
                                                                                                                                                                                                                                    Sequence 947 from Patent
AX215505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nogo gene expression
Patent: WO 0159103-A 51 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
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synthetic construct
artificial sequences.
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                                                                                                                 Blatt,L., Mcswiggen,J. and Chowrira,B.M. Method and reagent for the modulation and
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/mal_type="unassigned RNA"
/mb xref="taxon:32630"
/mb wref="Nucleic Acid"
organism="synthetic construct"
/mol_type="unassigned RNA"
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AX216978/c
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AX216404/c
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                                                                                         nogo gene expression
Patent: WO 0159103-A 2420 16-AUG-2001;
RIBOZYME DHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
                                                                                                                                                                                                                                           Sequence 2420 from Patent AX216978 AX216978.1 GI:15527039
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Blatt,L., Mcswiggen,J. and Chowrira,B.M.
Method and reagent for the modulation and diagnosis of cd20 and
nogo gene expression
Patent: WO 0159103-A 1846 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence (US);
MCSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
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AX216404
                                                                                                                                                       Blatt,L., Mcswiggen,J. and Chowrira,B.M. Method and reagent for the modulation and diagnosis of cd20 and
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                                    /organism="synthetic construct"
/mal_type="unassigned RNA"
/mb.xref="taxon:32630"
/mb.xref="Nucleic Acid"
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/mol_type="unassigned RNA"
/mb xref="taxon:32630"
/mbe="Nucleic Acid"
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/note="Nucleic Acid"
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AX262772
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AX216979/c
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16 ATGAACCGGAGG 27
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AX262772
AX262772.1
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Method and reagent for the modulation and dia
nogo gene expression
Patent: WO 0159103-A 2421 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
                                                                                                                                                                                   Kmiec,E.B., Gamper,H.B. and Rice,M.C. Targeted Chromosomal genomic alterations stranded oligonucleotides Patent: WO 0173002-A 163 04-OCT-2001; UNIVERSITY OF DELAWARE (US)
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                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; |
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                               Homo sapiens
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Conservative 0;
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/db xref="taxon:32630"
/noTe="Nucleic Acid"
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Pred. No. 9.1e+05;
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                                       RESULT 73
AX262801/c
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Sequence 164 from Patent
AX262773
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Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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llarity 100.0%; Pred. No. 9.1e+05;
Conservative 0; Mismatches 0;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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                                       Homo sapiens (human)
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Sequence 794 from Patent
AX499487
AX499487.1 GI:23381780
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Diagnosis and treatment of disease
Patent: WO 0244736-A 114 06-TUN-2002;
Molecular Skincare Limited (GB)
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             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Zhan, J.

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AX545291
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AX499493
AX499493.1 GI:23381786
                                                                                       Zhang, J., Gu, Y. and Nguyen, C.T.

Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase
Patent: Ep 1243660-A 804 25-SEP-2002;
                                                                                                                                                                        Homo sapiens (human)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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       /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 9.1e+05;
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AX545293
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165 CCACGTGGAATT 176
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12; Conserv
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Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase
Patent: EP 1243660-A 806 25-SEP-2002;
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                                                                                                                                                            Aeomica,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase
Patent: EP 1243660-A 805 25-8EP-2002;
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AX545292
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                      /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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100.0%; Pred. No. 9.1e+05;
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AX545294
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Sequence
AX545295
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Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase
Patent: Ep 1243660-A 807 25-SEP-2002;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Homo sapiens (human)
                           AX545296.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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807 from Patent EP1243660.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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ive 0; Mismatches 0;
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AX687580/c
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AX673640
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AX673640
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Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase
Patent: Ep 1243660-A 809 25-SEP-2002;
                                                                                 Homo sapiens
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Location/Qualifiers
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Sequences involved in phenomena of tumour
reversion, apoptosis and/or resistance to
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                          Shannon, M.,
                                                     Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                              Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent: WO 03004526-A 2085 16-JAN-2003;
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           non,M., Gu,Y. and Nguyen,C.T. human zinc-finger-containing
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2085 from Patent WO03004526.
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                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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100.0%; Pred. No. 9.1e+05;
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                                                     Craniata; Vertebrata; Catarrhini; Hominidae;
             proteins :
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             mdz3,
              mdz4, mdz7
                                                                  Euteleostomi;
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Aeomica,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; 
Mammalia; Butheria; Primates; Catarrhini; Hominidae;
                                                                                          Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins
                                                                                                                                                                                                      Sequence 793 from Patent AX688061
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                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                     Patent: EP 1281758-A 793
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ilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 9.1e+05;
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Pred. No. 9.1e+05;
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Best Local S
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Query Match 2.0%;
Best Local Similarity 100.0%;
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Best Local Similarity 100.0%;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                      Aeomica,
                                                                                                                                                                                       Patent: EP 1281758-A 795 05-FEB-2003;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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Aeomica, Inc. (US)
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llarity 100.0%;
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                                                                                  /organism="Homo sapiens"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 9.1e+05;
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Pred. No. 9.1e+05;
Score 12; DB 6; Li
Pred. No. 9.1e+05;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX688064
AX688064.1 GI:29410762
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 AX688066
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                             TTTACTGCTGGA 13
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Sequence 4428 from Patent
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/mol_type="unassigned DNA"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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12; Conservative 0; Mismatches 0;
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Sequence 4430 from Patent
AX691698
Patent: EP 1281758-A 4431 05-FEB-2003; 
Aeomica, Inc. (US)
                                  Shannon,M., Gu,Y. and Nguyen,C.T. Four human zinc-finger-containing proteins :
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                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Sequence 4432 from Patent
AX691700
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                                                                                                        Patent: EP 1281758-A 4433 05-FEB-2003;
                                                                                                                                       Shannon,M., Gu,Y. and Nguyen,C.T. Four human zinc-finger-containing proteins
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae,
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/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                          Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                      Location/Qualifiers
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Pred. No. 9.1e+05;
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Length 17;

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KEYWORDS
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AX724214
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AX724376/c
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Sequences involved in phenomena of tumour suppression, tumour sequences and their use as
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                                                                                                                       Patent: WO 03025176-A 2063 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                                         Telerman,A., Amson,R. and Tuijnder,M. sequences involved in phenomena of tumour suppression, tumour sequencies, apoptosis and/or virus resistance and their use as medicines.
                                                                                                                                                                                                                                                                                                                      Sequence 2063 from Patent AX724376
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                               Mus musculus (house mouse)
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Location/Qualifiers
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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                                                                 /organism="Mus musculus"
/mol_type="unassigned DNA"
/db_xref="taxon:10090"
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/mol_type="unassigned DNA"
/db_xref="taxon:10090"
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AX728700
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ORGANISM
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ACCESSION
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AX724877
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
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                                                                      12;
                                                                                                                                                                                                                                     Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumour
Sequences involved in phenomena of tumour suppression, tumour
                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (human)
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AX724877
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Conservative
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larity 100.0%;
Conservative (
                                                                                                                                                           /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:10090"
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AX734602
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ORGANISM
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TITLE
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   AX734602 17 bp 1
Sequence 192 from Patent W003025177
AX734602
AX734602.1 GI:30513879
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Sequences involved in phenomena of tumour suppression,
reversion, apoptosis and/or virus resistance and their
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Sequence 2060 from Patent WO03025175.
AX730426
AX730426.1 GI:30509769
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Sequence 3542 from Patent
AX731908
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                   n 2.0%; Score 12; DB 6; Length 17;
Similarity 100.0%; Pred. No. 9.1e+05;
12; Conservative 0; Mismatches 0; Indels
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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AX736368/c
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AX734928/c
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Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                              AX736368
Sequence 1958 from Patent
AX736368
AX736368.1 GI:30515645
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Patent: WO 03025177-A 518 27-MAR-2003;
Molecular Engines Laboratories (FR)
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AX734928
AX734928.1 GI:30514205
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Patent: WO 03025177-A 192 27-MAR-2003;
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Location/Qualifiers
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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larity 100.0%;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Sequences involved in phenomena of tumour suppression,
reversion, apoptosis and/or resistance to viruses and t
thereof as medicaments
Patent: WO 03025177-A 5159 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and the use thereof as medicaments
Patent: WO 03025177-A 1958 27-MAR-2003;
Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in tumoral suppression, tumoral reversion, apoptosis and/or viral resistance phenomena and their use as medicines
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                                                                                                                Homo sapiens (human)
                                                                                                                                                                                 Sequence 1057 from Patent AX757736
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
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Sequence
AX762288
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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                            Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in tumoral suppression, tumoral reversion.
                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                            Homo sapiens (human)
                                                                                                                                                                                                                                                AX762288.1 GI:32256904
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent: WO 03040369-A 3518 15-MAY-2003;
Molecular Engines Laboratories (FR)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                               Patent: WO 03040369-A 5609 15-MAY-2003;
                                                                                               apoptosis and/or viral resistance phenomena and their use
                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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5609 from Patent WO03040369.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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organism="Homo sapiens"/
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Pred. No. 9.1e+05;
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Journal Ja; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

1 (bases 1 to 17)

Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A.

TITLE molecule participating in vasculogenic response

Patent: JP 2002509721-A 1790 02-APR-2002;

OMMENT OS Homo sapiens (human)

PN JP 2002509721-A/1790

PD 02-APR-2002

PP 27-MAR-1998 US

PI PAMELA A PANYO

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PC

Concerniation Conditions concerning

PN JP 2002509721-A/1790

PR 27-MAR-1999 US

PR 27-MAR-1999 US

PI PAMELA A PANYO

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Concerniation Con
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BD200907
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BD198764/c
                 VERSION
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17 bp RNA linear PAT 17-JUL-200 Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response.

BD200907
BD200907.1 GI:33010677
BD2009509721-A/3933.
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PC A61P35
C12N5/00
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/mol_type="genomic RNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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100.0%; Pred. No. 9.1e+05;
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1 (bases 1 to 17)

Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A. Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response Patent: JP 2002509721-A 3933 02-APR-2002; RIBOZYME PHARMACEUTICALS INC

OS Homo sapiens (human)
PN JP 2002509721-A/3933
PD 02-APR-2002
                                                                                                                                                             artificial sequences.

artificial sequences.

1 (bases 1 to 18)

2abeau,M. and Vos,P.

Selective restriction fragment amplification : a general method for DNA fingerprinting

Patent: EP 0534858-A 43 31-MAR-1993;
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PC A61P35
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27-MAR-1998 US 60/079678
PAMELA A PAVCO, ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT,
JAMES A MCSWIGGEN
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Key Location/Qualifiers
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/mol_type="genomic RNA"
/db_xref="taxon:9606"
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Location/Qualifiers
                                                                                 /organism="synthetic construct"
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AR085592
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Sequence 13
AR092798
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Monia,B.P. and Cowsert,L.M.
Antisense modulation of MEK2 expression
Patent: US 5959097-A 17 28-SEP-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                AR085592
Sequence 28 from patent US
AR085592
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                                                                                                                                                                                                l Similarity
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Unclassified.
1 (bases 1 to 18)
Cowsert, L.M.
                                                                 AR092798.1 GI:10019550
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Cowsert, L.M.
                                                                                                                                                                                                                                                                                                                                           Unclassified.
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                                 Unknown.
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                                                                                                                                                                                                                                                                                                                                                    synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 18)
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AR096294
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Patent: JP 2002531130-A 8 24-SEP-2002;
CHILDREN'S HOSPITAL AND REGIONAL MEDICAL CENTER
OS Artificial Sequence
PN JP 2002531130-A/8
PD 24-SEP-2002
PP 08-DEC-1999 JP 2000586917
PR 08-DEC-1998 US 60/111493
PI PHILLIP I TARR
PC C12N15/09,C12N15/09,C07K16/12,C12N1/15,C12N1/19,C12N1/21
,C12N5/10,C12N9/04,
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/mol_type="unassigned
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Query Match
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Sequence
127949
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Apple,R.J., Erlich,H.A., Griffith,R.L. and Scharf,S.
Methods and reagents for HLA DRbeta DNA typing
Patent: US 5567809-A 121 22-OCT-1996;
Location/Qualifiers
                                               1 (bases 1 to 18)
Apple,R.J., Erlich,H.A., Griffith,R.L. a
Methods and reagents for HLA DRbeta DNA
Patent: US 5567809-A 158 22-OCT-1996;
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C12P21/08,
PC (C12N9/04, C12R1:19), (C12Q1/68, C12R1:19), C12N15/00, C12N5/00
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/mol_type="unassigned DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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             organism="unknown"
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Sequence 17
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Kluyveromyces.
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                                                                                                                                                                                                       fungal pathogens
Patent: WO 0073499-A 8 07-DEC-2000;
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Morin,G.B., Funk,W.D. and Piatyszek,M.
Second mammalian tankyrase
Patent: US 6599728-A 17 29-JUL-2003;
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                                                                                                            /organism="Kluyveromyces marxianus"
/mol_type="unassigned DNA"
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REFERENCE AUTHORS TITLE

RESULT 119 127949 LOCUS

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synthetic synthetic
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Location/Qualifiers
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Location/Qualifiers
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65 from Patent EP1184468.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Sample oligonucleotide"
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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Sequence 1 from Patent EP1188475.
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Screening method for gene variation
Patent: EP 1184467-A 65 06-MAR-2002;
CANON KABUSHIKI KAISHA (JP)
Location/Qualifiers
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                    Okamoto,T., Yamamoto,N., Watanabe,H. and Suzuki,T. Method for making probe support and apparatus used Patent: EP 1188475-A 1 20-MAR-2002; CANON KABUSHIKI KAISHA (JP)
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Sequence
AX453818
                                                                                                                                                                               Okamoto,T., Yamamoto,N. and Suzuki,T.
Terminal labeled probe array and method
Patent: EP 1213361-A 65 12-JUN-2002;
CANON KABUSHIKI KAISHA (JP)
                                                                                                                                                                                                                                                                                                                           Sequence 65 from Patent EP1213361.
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Location/Qualifiers
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               2.0%; Sc
llarity 100.0%; P
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42 from Patent EP1213361.
                                                                                      /organism="synthetic construct"
/mol_type="unassigned DNA"
/mb xref="taxon:32630"
/note="Synthesized"
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/db_xref="taxon:32630"
/noTe="Base Oligonucleotide for preparation
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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9.1e+05;
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9.1e+05;
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CE 1 (bases 1 to 18)

RS Okamoto,H., Yamamoto,N. and Suzuki,T.

Probe-coupling substrate, process for producing the same,
probe-array, method for detecting target substance, method for
specifying base sequence of single-stranded nucleic acid in sample
and method for quantitating the target substance in the sample
and method for quantitating the target substance in the sample
Patent: JP 2000270896-A, 1 03-OCT-2000;

RTLIFICIAL Sequence
PN JP 2000270896-A/1
PD 03-OCT-2000

PF 28-JAN-1999 JP 1999019915
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12; Conserv
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BD000011.1 GI:18623090
JP 2000270896-A/1.
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                                                                                                    HISASHI OKAMOTO, NOBUKO YAMAMOTO, TOMOHIRO SUZUKI
1/68, C12M1/00, C12N15/09, G01N33/566, C12N15/00 CC
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                                                                                        Key
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34 from Patent WO0236751.
                                         /organism='Artificial Location/Qualifiers
/organism="synthetic construct"
/mol_type="genomic DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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RESULT 132
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OS Artificial Sequence
PN JP 2000270896-A/43
PD 03-OCT-2000
PP 28-JAN-1999 JP 1999019915
PR 11SASHI OKAMOTO,NOBUKO YAMAMOTO,TOMOHIRO SUZUKI PC C12Q1/68,C12M1/00,C12N15/09,G01M33/566,C12N15/00 CC
FH Key 1. 18
FT source /organism='Artificial Sequence'.
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synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 18)
                                                           BD010932.1 GI:18639305
JP 2001061486-A/42.
                                                                                           fingerprint method Selegeneral DNA fingerprint BD010932
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1 (bases 1 to 18)
0kamoto, H., Yamamoto, N. and Suzuki,
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BD000053.1 GI:18623132
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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100.0%; Pred. No. 9.1e+05;
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Best Local S
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Best Local Similarity 100.0%; F
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OS Artifio
PN JP 201
PD 13-MAR
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PR MARK S;
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16 ATGAACCGGAGG
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OS Artif
PN JP 20
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(bases 1 to 18)

Yamamoto,N., Okamoto,T., Suzuki,T. and Tanaka,S.
Method for screening mutated gene
Patent: JP 2002071687-A 42 12-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic construct
synthetic construct
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Method for screening mutated gene.
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JF 2002071687-A/42
12-MAR-2002
112-MAR-2000
31-AUG-2000 JF 2000263396
NOBUKO YAMAMOTO,TADASHI OKAMOTO,TOMOHIRO SUZUKI,SHINYA TANAKA
ROBUKO YAMAMOTO,TADASHI OKAMOTO,TOMOHIRO SUZUKI,SHINYA TANAKA
GO1N33/53,C12M1/00,C12N15/09,C12Q1/68,G01N31/22,G01N33/566, PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUL-2000 JP 2000224187
24-SEP-1991 GB 91402542.4
MARK SZABO, PIETER FOSS
C12N15/09, C12Q1/68, C12N15/00
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                                                                                                                                                                                             /organism='Artificial Location/Qualifiers
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                                                                                                                   /organism="synthetic construct
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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|mol_type="genomic DNA"
|db_xref="taxon:32630"
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                                                    Score 12; DB 6; L
Pred. No. 9.1e+05;
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BD135742/c
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BD135697/c
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                                                    Patent:
CANON IN
OS Art
PN JP
PD 05-
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  probe for mutated
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OS Artifi
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PP 31-AUG
PI NOBUKO
PC C12N15
G01N33/566,
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synthetic construct
artificial sequences.
                                                                                                                                              Yamamoto,N., Okamoto,T., Suzuki,T. and Shimizu,A.
Method for detecting subjective component in specimen sample, and
substrate for detection used therefor
Patent: JP 2002065274-A 46 05-MAR-2002;
                                                                                                                                                                                                                                                                                     substrate for detection BD135742
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                                      G01N33/566
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1 (bases 1 to 18)
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JP 2002065274-A/46.
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                          G01N35/02,G01N35/10,G01N37/00,C12N15/00,G01N35/06 CC
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                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-AUG-2000 JP 2000263395
NOBUKO YAMAMOTO,TADASHI OKAMOTO,TOMOHIRO SUZUKI,AKIRA SHIMIZU NOBUKO (12M1/00,C12M1/40,C12Q1/68,G01N31/22,G01N33/53, PC
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JP 2002065274-A/1
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 or hybridizing with gene encoding p53; named
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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100.0%; Pred. No.
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BD161031
BD161031.1 GI:27866789
JP 2002153284-A/65.
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artificial sequences.

1 (bases 1 to 18)

0 kamoto, T., Yamamoto, N. and Suzuki, T.
Terminal-labeled probe-array and method method for evaluating target mass using Patent: JP 2002153284-A 42 28-MAY-2002; CANON INC.
BD161031 18 bp DNA linear PAT 1
Terminal-labeled probe-array and method for preparing it,
method for evaluating target mass using the same.
                                                                                                                                                                                                                                                                                                                                                                                                                                             PF 24-NOV-2000 JP 2000357446
PI TADASHI OKAMOTO, NOBUKO YAMAMOTO, TOMOHIRO SUZUKI PC
C12N15/09, C12Q1/68, G01N31/22, G01N33/53, G01N33/566, G01N37/00,
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synthetic construct

synthetic construct
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BD161008.1 GI:27866766
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Terminal-labeled probe-array
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JP 2002153284-A/42
28-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                             Description of Artificial Sequence:Synthesized FH
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Location/Qualifiers
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/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                     Best Local Similarity Matches 12; Conserv
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PI TADASHI OKAMOTO, NOBUKO YAMAMOTO, TOMOHIRO SUZUKI PC C12N15/09, C12Q1/68, G01N31/22, G01N33/53, G01N33/566, G01N37/00,
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1 (bases 1 to 18)

Okamoto, T., Yamamoto, N. and Suzuki, T.

Okamoto Terminal - labeled probe-array and method method for evaluating target mass using Patent: JP 2002153284-A 65 28-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         synthetic construct
synthetic construct
artificial sequences.
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28-MAY-2002
                   ilarity 100.0%; P
Conservative 0;
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12-DEC-2000 JP 2000377349
TOMOHIRO SUZUKI
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                       /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                        2.0%; Score 12;
100.0%; Pred. No.
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BD162059
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PD 25-JI
PF 12-DE
PF TOMOD
PC C120J
PC C12NJ
PC C12NI
P
                                                                                                                    Yamamoto,N., Okamoto,T. and Suzuki,T.

A method of analyzing a base sequence of a nucleic acid Patent: WO 0233068-A 42 25-APR-2002;
CANON KK,NOBUKO YAMAWOTO,TADASHI OKAMOTO,TOMOHIRO SUZUKI OS Artificial Sequence
N WO 0233068-A/42
PM 0233068-A/42
PD 18-OCT-2000 WO 2000JP007244
PT 18-OCT-2000 WO 2000JP007244
PT 18-OCT-2000 WO 2000JP007244
PT 18-OCT-2000 WO 2000JP007244
PT 18-OCT-2000 WO 2000JP007244
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                     C12N15/09, C12Q1/68, G01N33/586, G01N33/53
CC Sample origonucleotide
FH Key Location/Qualifier
FT source /organism='Artific:
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WO 0233068-A/42.
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TOMOHIRO SUZUKI
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/organism='Artificial Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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BD174965
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Liquid discharge device for production of probe carrier, device for production of probe carrier using the liquid discharge device, and process for producing the probe carrier patent: JP 2002257694-A 1 11-SEP-2002;

CANON INC

OS Artificial Sequence
PN JP 2002257694-A/1

PD 11-SEP-2002
PF 28-FEB-2001
PD 11-SEP-2002
PF 28-FEB-2001
PG 301N1/00, HAIJZ/05, CIZM1/00, CIZN15/09, G01N33/53, G01N33/566, PC G01N35/10, G01N35/10,
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                                                            and device used for thi
BD174965
BD174965.1 GI:29120659
JP 2002257836-A/1.
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Process for producing probe carrier using and device used for this process.
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BD174790.1 GI:29120482
JP 2002257694-A/1.
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                   synthetic construct synthetic construct
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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9.1e+05;
hes 0;
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discharge device
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Best Local S
Matches 12
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Patent: JP 2002257836-A 1 11-SEP-2002;
CANON INC
OS Artificial Sequence
PN JP 2002257836-A/1
PD 11-SEP-2002
PF 28-FEB-2001 JP 2001055971
PI TADASHI CKAMOTO
PC G01N35/10,B41J2/01,B41J2/05,C12M1/0
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PC B41J3
CC Base
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Kaneko, M. and Watanabe, H.
A method of preparing a probe array
Patent: JP 2002253251-A 1 10-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                synthetic construct synthetic construct
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JP 2002253251-A/1.
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                                                                                                 Similarity
                                      ATGAACCGGAGG 27
                                                                                                                                                                                                                                                                                                                                                      P 28-FEB-2001 JP 2001055972
I MINEO KANEKO, HIDENORI WATANABE
CCLEN15/09, C12M1/00, C010133/53, C010133/566, G01N37/00, C12N15/00 CC
Base Oligonucleotide for preparation of a probe FH Key
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ATGAACCGGAGG
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                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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ilarity 100.0%;
Conservative
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Base Oligonucleotide for preparation
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G01N35/10,B41J2/01,B41J2/05,C12M1/00,C12N15/09,G01N37/00,
                                                                                Conservative
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                                                                                                                                                                                   /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Location/Qualifiers
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mol_type="genomic DNA"
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9.1e+05;
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                                source
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PN JP 2002306166-A/42
PD 22-0CT-2002
PF 31-AUG-2000 JP 2000263506
PI NOBUKO YAMAMOTO, HISASHI OKAMOTO, TOMOHIRO SU:
C12N15/09,C12Q1/68//C12M1/00,C12N15/00
CC Sample origonucleotide
FH Key Location/Qualifiers
FT source 1.18
FT source /organism='Artificial Sequen
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BD176986.1 GI:30014245
JP 2002306166-A/42.
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                                                            PC G01N33/53,G01N37/00//C12M1/00,C12N15/09,C12N15/00 CC Oligonuclectide for preparation of a probe FH Key Location/Qualifiers
FT Source 1. 18
FT fource /organism='Artificial Sequence'.
                                                                                                                                                                                                                                                    artificial sequences.

1 (bases 1 to 18)

Watanabe,H., Okamoto,T., Yamamoto,N. and Suzuki,T.

A method of preparing a probe array and a device used therefor Patent: JP 2002318232-A 1 31-OCT-2002;
                                                                                                                                                                                                                                                                                                                                                                                                 A method of preparing a probe array and a device used therefor BD177274
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                                                                                                                                                      Artificial Sequence
JP 2003318232-A/1
31-OCT-2002
18-SEP-2001 JP 2001283190
HIDENORI WATANABE,TADASHI OKAMOTO,NOBUKO YAMAMOTO,TOMOHIRO
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Location/Qualifiers
                                       /organism='Artificial
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
/organism="synthetic construct"
/mol_type="genomic DNA"
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9.1e+05;
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AUTHORS
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BD187512/c
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KEYWORDS
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1 (bases 1 to 18)

Okamura, N., Okamoto, T. and Kameyama, M.

Probe carrier, Method and Apparatus for producing Probe carrier pratent: JP 2003014773-A 1 15-JAN-2003;
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BD187511.1 GI:32997250
JP 2003014773-A/1.
                                                                                                        artificial sequences.

1 (bases 1 to 18)

Okamura, N., Okamoto, T. and Kameyama, M.

Probe carrier, Method and Apparatus for producing Probe carrier

Patent: JP 2003014773-A 2 15-JAN-2003;
                                                                                                                                                                                         synthetic construct synthetic construct
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                                                                                                                                                                                                                                                       Probe carrier, Method and BD187512
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oligonucleotide used as
                                                                                             CANON IN
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JP 2003014773-A/1
15-JAN-2003
                                             Artificial Sequence JP 2003014773-A/2 15-JAN-2003
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                28-MAR-2002 JP 2002093024 nobuyuki okamura, tadashi o
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/mol_type="genomic DNA"
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         Narwa,R. and Roques,P.

Nucleic acid fragments derived form the HIV-1 genome, corresponding fragments and their application as reactives for risk evaluation of HIV-1 mother-foetal transmission gatent: EP 0743364-A 34 20-NOV-1996; Patent: EP 0743364-A 34 20-NOV-1996; COMMISSARIAT ENERGIE ATOMIQUE (FR)

COMMISSARIAT ENERGIE ATOMIQUE (FR)

Other publication FR 2734281 961122.
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Sequence 34 from Patent EP0743364.
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Nucleic acid fragments derived form the HIV-1 genome, corresponding fragments and their application as reactives for risk evaluation of HIV-1 mother-foetal transmission patent: EP 0743364-A 33 20-NOV-1996, COMMISSAINT EMERGIE ATOMIQUE (FR)

Other publication FR 2734281 961122.
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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Nucleic acid fragments derived form the HIV-1 genome, corresponding fragments and their application as reactives for risk evaluation of HIV-1 mother-foetal transmission Patent: EP 0743364-A 35 20-NOV-1996; COMMISSARIAT ENERGIE ATOMIQUE (FR)

Other publication FR 2734281 961122.

Location/Qualifiers
                                                                                                                                          1 (bases 1 to 19)
Pallisgaard,N. and Hokland,P.
PETECTION OF CHROMOSOMAL ABNORMALITIES
PACKETHE WO 9824928-A 54 11-UN-1998;
PALLISGAARD NIELS (DK); HOKLAND PETER
LOCATION/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
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Sequence 14 from Patent WC0015788
AX023306
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Cohen,D., Chumakov,I. and Blumenfeld,M.
Biallelic markers for use in constructing a
disequilibrium map of the human genome
Patent: US 6537751-A 11302 25-MAR-2003;
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12; Conserv
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AR299567
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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/mol_type="genomic
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                                       synthetic construct synthetic construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Agrobacterium tumefaciens (Rhizobium radiobacter)
Agrobacterium tumefaciens
Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
Rhizobiaceae; Rhizobium/Agrobacterium group; Agrobacterium.
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CENTRE NATIONAL DE LA RECHERCHE SCIENTIFIQUE
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  Roden, R. and Naora, H.
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/mol_type="unassigned DNA"
/db_xref="taxon:358"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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AX763550
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AX763549
AX763549.1 GI:32257984
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Patent: WO 0168853-A 8 20-SEP-2001;
The Johns Hopkins UNiversity School
Location/Qualifiers
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                                              Harel-Bellan, A., Ait-Si, S., Cabon-Georget, F.,
                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                              Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                       Inhibitor oligonucleotides and their use for
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 WO 03040366-A 63
                                                                                                                                                        63 from Patent W003040366
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:32630"
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AX763551
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Rujan, T. and Schmitt, A. Method and nucleic acids for the analysis proliferative disorder Patent: EP 1340818-A 417 03-SEP-2003; Epigenomics AG (DE)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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Location/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Location/Qualifiers

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AX826165/c
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ORGANISM
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BD023309/c
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E 1 (bases 1 to 19)

S Parisgard.N. and Hokurando, P.

S Method for detecting abnormality in chromosome Patent: JP 2001505428-A 54 24-APR-2001;

NEILLS PARISGARD PN JP 2001505428-A/54

PD 24-APR-2001

PF 08-DEC-1997 JP 1998525090

PF 08-DEC-1997 JP 1998525090

PF NEILLS PARISGARD, PATER HOKURANDO C12N15/09,C12Q1/6B,G01N33/50,C12N15/00

CC Strandedness: Single;

CC Topology: Linear;
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Method fo
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JF 2001505428-A/54.
Homo sapiens (human)
Homo sapiens
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Patent: WO 03072821-A 417 04-SEP-2003;
Epigenomics AG (DE)
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Sequence 417 from Patent
AX826165
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                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                       BD023309
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for detecting abnormality in
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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100.0%; Pred. No. 9.1e+05;
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100.0%; Fred. No. 9.1e+05;
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DEFINITION
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AR085567
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Sequence 35 from Patent WO9854360.
A82447
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Sequence
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                                                                                               Antisense modulation of G-alpha-13 expression Patent: US 5981732-A 3 09-NOV-1999; Location/Qualifiers
                                                                                                                                   1 (bases 1 to 20)
Cowsert, L.M.
                                                                                                                                                                        Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unclassified.

(bases 1 to 20)

Plastow,G.S. and Wales,R.

Plastow,G.S. ANALYZING ANIMAL PRODUCTS

METHODS FOR ANALYZING ANIMAL PRODUCTS

PATENT: MO 9854360-A 35 03-DEC-1998;

PLASTOW GRAHAM STUART (GS); WALES RICHARD

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             unidentified
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                                                                                                                                                                                      Unknown.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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                                                           /organism≈"unknown"
/mol_type="unassigned DNA"
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                                                                                                                                                                                                              GI:10012334
2.0%; Score 12; DB 6; L
100.0%; Pred. No. 9.1e+05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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Pred. No. 9.1e+05;
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5981732.
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9.1e+05;
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                        Length 20
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                                                                                                                                                                                                                                                  linear
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Conservative

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PAT 16-JUN-2001

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KEYWORDS
SOURCE
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AUTHORS
TITLE
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AR099487
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AUTHORS
TITLE
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AR097384
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ACCESSION
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AR097385/c
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                      AR099487
Sequence 14
AR099487
AR099487.1
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Sequence
AR097385
                                                                                                                                                                                                                                                                            1 (bases 1 to 20)
Diamandis, E., Dunn, J.M. and Stevens, J.K.
Method, reagents and kit for diagnosis a:
p53 mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
AR097384
                                                                                                                                                            12;
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Diamandis, E., Dunn, J.M. and Stevens, J.:
Method, reagents and kit for diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9
  Unknown
                                                                                                                                                                                                                                                     Patent: US 6071726-A 9 06-JUN-2000;
Location/Qualifiers
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12; Conserv
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Location/Qualifiers
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                                                                                                                                                                                                                   /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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                                            from patent US 6077833.
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6071726.
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9.1e+05;
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AUTHORS
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AUTHORS
TITLE
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VERSION
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AR143130/c
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ACCESSION
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AR137400
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Query Match
Best Local Similarity
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Unclassified.

1 (bases 1 to 20)

1 (bases 1 to 20)

Berg, T., Tollersrud, O. Kristien. and Nilssen, O.

Genetic test for .alpha.-mannosidosis

Patent: US 6197507-A 15 06-MAR-2001;

Location/Qualifiers
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Sequence
AR143130
AR143130.
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AR137400
                                                                            1 (bases 1 to 20)
Dean,N.M. and Marcusson,E.G.
Antisense inhibition of Fas mediated
Patent: US 6204055-A 21 20-MAR-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                        l Similarity
12; Conserv
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Bennett, C. Frank. and Vickers, T.A.
Oligonucleotide compositions and methods for the modulation expression of B7 protein
Patent: US 6077833-A 14 20-JUN-2000;
                                                                                                                                                 Unknown
                                                                                                                                                           Unknown.
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(larity 100.0%;
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                                                                                                                                                                                                        21
                                           /organism="unknown"
/mol_type="unassigned DNA'
                                                                                                                                                                                                                                                                                                                                                                              /mol_type="unassigned
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/mol_type="unassigned
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 2.0%;
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0; Mismatches
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Pred. No.
 Score 12;
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9.1e+05;
 DB 6; Length
9.1e+05;
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PAT 08-AUG-2001

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BD230144/c
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AUTHORS
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AR178768
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DEFINITION
ACCESSION
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AR178952
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AR178768
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Sequence
AR178952
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Bennett, C. Frank. and Vickers, T.A.
Oligonucleotide compositions and methods
expression of B7 protein
Patent: US 6319906-A 14 20-NOV-2001;
                                                                                                                                                                                                                                                                               Unclassified.

(bases 1 to 20)

Bennett, C.Frank: and Vickers, T.A.

Bennett of Errank: and vickers, T.A.

Oligonucleotide compositions and methods expression of B7 protein

Patent: US 6319906-A 198 20-NOV-2001;
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 Total genome radiation hybrid map of canine identification of interesting genes. BD230144
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                                                                                                                                                                                                                                                                                                                                                                                                                                  198 from patent
                                                                                                                                                                                                                           /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned
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100.0%; Pr
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0; Mismatches
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Pred. No. 9.1e+05;
0; Mismatches 0;
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OS Canis familiaris (dog)
PN JP 2002530091-A/13
PD 17-SEP-2002
PF 15-NOV-1999 UP 2000582596
PF 13-NOV-1999 UP 2000582596
PF 13-NOV-1998 US 60/108193
PJ FRANCIS GALLBERT, CATHERINE ANDRE
PC C12N15/09,C12Q1/68,C12N15/00
CC Ren01023
FM Key Location/Qualifiers
FM Key 1.20
FT Source 1.20
FT Source 1.20
                                                                                                                                                                     artificial sequences.

1 (bases 1 to 20)

Dean, N.M. and Marcusson, E.G.
Antisense modulation of FAS mediated signaling
Patent: JP 2002540812-A 20 03-DEC-2002;

ISIS PHARMACEUTICALS INC
OS Artificial Sequence
PN JP 2002540812-A/20
PD 03-DEC-2002
PD 03-DEC-2000 JP 200610483
PR 12-APR-1999 US 09/290640
PI NICHOLAS M DEAN ERIC G MARCUSSON
PI NICHOLAS M DEAN ERIC G MARCUSSON
PI NICHOLAS M DEAN ERIC G MARCUSSON
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PC A61P
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Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
1 (bases 1 to 20)
Galibert,F. and Andre,C.
                                                                                                                                                                                                                                                                                                                                                                                         Antisense modulation of FAS m
BD249305
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                                                      SSEFF
                                                                                                                                                                                                                                                                                                                                    synthetic construct
synthetic construct
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Canis familiaris
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JP 2002530091-A/13.
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                                                                                                                                  A61P1/16, A61P29/00, A61P35/00, A61P37/00, A61P43/00//C12N5/06, PC
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                                                                               Synthetic Sequence
Loc
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                                                                                                                                                            C12N15/09,A61K31/7088,A61K31/7115,A61K31/712,A61K31/7125, PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism='Canis familiaris (dog)'.
                                     /organism='Artificial
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Canis familiaris"
/mol_type="genomic DNA"
/db_xref="taxon:9615"
/organism="synthetic construct"
/mol_type="genomic DNA"
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I25689
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                                                                                                                                                                                                                                                                                                                                                                                    Patent: JP 1993337000-A 31 21-DEC-1993;
CHAYAMA KAZDAKI

OS Artificial gene
OC Artificial sequence; Genes.
OS Hepatitis C virus
PN JP 1993337000-A/31
PD 21-DEC-1993
PP 04-JUN-1992 JP 1992168226
PI CHAYAMA KAZDAKI, KUMADA HIROMITSU
PC C12Q1/68,C12N15/10,C12N15/11,C12Q1/70
CC strandedness: Single;
CC topology: Linear;
CC hypothetical: No.
CC anti-sense: No.
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                        1 (bases 1 to 20)
Diamandis, D. Dunn, J.M. and Stevens, J.K.
Method, reagents and kit for diagnosis and targeted screening
P53 mutations
                                                                                                                                                                                                                     œ
                                                                                                                                   Sequence 8 from patent US 125689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    artificial sequences.

1 (bases 1 to 20)

Chayama, K. and Kumada, H.
METHOD FOR EXAMINING C T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E06107
Oligonucleotide specific
            Patent: US 5552283-A 8 03-SEP-1996;
                                                                                 Unknown
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synthetic construct
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JP 1993337000-A/31.
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ilarity 100.0%;
Conservative (
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CHAYAMA KAZUAKI, KUMADA HIROMITSU
C12Q1/68,C12N15/10,C12N15/11,C12Q1/70;
strandedness: Single;
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                           /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                       GI:1605559
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Query Match
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180 CTTCCTCCGCTA 191
                                                                                                                                             1 (bases 1 to 20)
Szostak,J.W. and Huizenga,D.E.
DNA aptamers and castalysts that bind adenosine adenosine-5'-phosphates and methods for isolati Patent: US 5631146-A 123 20-MAY-1997;
                                                                                                                                                                                                                                                                                Sequence 123 from patent I43305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 20)
Diamandis, E., Dunn, J.M. and Stevens, J.K.
Method, reagents and kit for diagnosis and targeted
P53 mutations
                                                                                                                                                                                                               Unclassified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent: US 5552283-A 9 03-SEP-1996;
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                                      Similarity
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ilarity 100.0%;
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                                                                                                                               Location/Qualifiers
                                                                                        /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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US 5631146.
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3. 9.1e+05;
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.9.1e+05;
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Pouzyrev, A. Timofeyevich. and Riddle, D. Lee.
Method for construction of cDNA libraries
corresponding to rare mRNA
Patent: US 6372436-A 1 16-APR-2002;
Location/Qualifiers
                                                                                                 Sequence 6 1
AR281883
                                                                                                                                                                                                                       12;
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Unclassified.

1 (bases 1 to 20)

Warenius,H.M. and Seabra,L.A.

Methods for determining chemosensitivity of cancer cells based upon expression of negative and positive signal transduction factors
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 172 from patent AR265991 AR265991.1 GI:29694837
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ilarity 100.0%;
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/mol_type="unassigned
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/mol_type="genomic
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Sequence 11
AR299997
AR299997.1
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Griffals,R., Hoiseth,S.K., Zagursky,R.J.,
Sankaran,B. and Fletcher,L.D.
Chlamydla pneumoniae polynucleotides and
Patent: US 6559294-A 1338 06-MAY-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 20)
Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 11732 25-MAR-2003;
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Location/Qualifiers
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Sequence 1338 from patent
AR310801
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/mol_type="genomic
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/mol_type="genomic DNA"
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AR337679
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AR437041/c
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AR437041
AR437041.1
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Dean, N.M., Marcusson, E.G. and Wyatt, J.

Antisense modulation of Fas mediated s

Patent: US 6633133-A 21 25-NOV-2003;

Location/Qualifiers
            Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 from patent US 6653133. AR432224
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Wright, J.A., Young, A.H. and Lee, Y.S.
Oligonucleotide sequences complementary to thioredoxin or
Chioredoxin reductase genes and methods of using same to o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14 from patent US 6566514.
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Patent: WO 9942839-A 6 26-AUG-1999;
THERYTE LIMITED (GB); WARENIUS HILMAR (GB)
Location/Qualifiers
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Bennett,C.F. and Watt,A.T.
Antisense inhibition of src-c expression
Patent: US 6656732-A 93 02-DEC-2003;
Location/Qualifiers
                                                                                  Seabra, L.A. and Warenius, H.
Treating cancer
Patent: WO 9942834-A 6 26-AUG-1999;
SEABRA LAURENCE ANTHONY (GB); THERYTE LIMITED
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llarity 100.0%;
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 /organism="synthetic construct"
/mol type="unassigned DNA"
/mol type="taxon:32630"
/mbe="paymer"
                                                           Location/Qualifiers
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/note="PRIMER"
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from Patent WO9942834.
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Pred. No. 9.1e+05;
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Patent: WO 9942828-A 6 26-AUG-1999;
THERYTE LIMITED (GB); WARENTUS HILMAR MEEK
Location/Qualifiers
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Treating cancer
Patent: WO 9942821-A 6 26-AUG-1999;
SEABRA LAURENCE ANTHONY (GB); THERYTE LIMITED (GB); WARENIUS HILMAR
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db xref="taxon:32630"
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                                      2.0%; Score 12; DB 6; Le
100.0%; Pred. No. 9.1e+05;
o. Mismatches 0;
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Sequence 5413 from Patent WO0179548.
AX293651
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Sequence 6 from Patent WO9942090.
AX019035
AX019035.1 GI:10043116
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THERYTE LIMITED (GB); WARENIUS HILMAR MEEK (GB)
Location/Qualifiers
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100.0%; Pred. No. 9.1e+05;
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AX297041/c
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Sequence 8803 from Pate:
AX297041
AX297041.1 GI:17058732
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Barany, F., Zirvi, M., Gerry, N.P., Favis, R.
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ilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                            /note="Hypothetical Probe Sequence"
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AX459997
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AX301845
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Sequence 6 from Patent WO0185917.
AX301845
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                                           Microarray method of genotyping multiple samples at Patent: WO 0203849-A 8 17-JAN-2002;
Telechem Intnational, Inc. (US)
Location/Qualifiers
                                                                                                                                                                 Homo sapiens
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Sequence 8 from Patent WO0203849.
AX459997
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                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/organism="Homo sapiens"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="primer"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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AX613711
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AX613711
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Sequence 265 from Patent EP1288296.
AX710965
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                                                                                                                      Draper,K.G., Mcswiggen,J.A., Holecek,J.J., Dudycz,L.W., Macejak,D.G. and Mamone,J.A. Method and reagent for inhibiting HBV viral replication Patent: EP 1288296-A 265 05-MAR-2003; RIBOZYME PHARMACEUTICALS, INC. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent: WO 0207
OGHAM GmbH (DE)
                                                                                                                                                                                                                                  Human herpesvirus 5
Human herpesvirus 5
Viruses; dsDNA viruses, no RNA stage;
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Mammalia; Eutheria; Primates;
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              Similarity
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llarity 100.0%; Pred. No. 9.1e+05;
Conservative 0; Mismatches 0;
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2.0%; Score 12; DB 6; Locality 100.0%; Pred. No. 9.1e+05; Conservative 0; Mismatches 0;
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4736 from Patent WO02072882.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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                                                                /organism="Human herpesvirus
/mol_type="unassigned RNA"
/db_xref="taxon:10359"
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PC C12N5/00,
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15 OCT 1992 US
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JP 2000342285-A/266.
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Method and reagent for inhibiting viral replication.
BD001535
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JP 2000342286-A/266.
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11-MAY-1992 US 07/882689,14-MAY-1992 US 07/882713,14-MAY-1992 US 07/882713,14-MAY-1992 US 07/882892 J4-MAY-1992 US 07/882892 J4-MAY-1992 US 07/882892 J4-MAY-1992 US 07/882896,14-MAY-1992 US 07/882899,14-MAY-1992 US 07/882899,14-MAY-1992 US 07/882899,14-MAY-1992 US 07/882891 US 07/882899,14-MAY-1992 US 07/882891 US 07/882899,14-MAY-1992 US 07/88249,14-MAY-1992 US 07/88249,14-MAY-1992 US 07/88291,14-MAY-1992 US 07/88349,14-MAY-1992 US 07/88349,14-MAY-1992 US 07/88349,14-MAY-1992 US 07/88349,14-MAY-1992 US 07/884436,14-MAY-1992 US 07/884436,14-MAY-1992 US 07/8839,26-AUG-1992 US 07/963322,07-DEC-1992 US 07/963322,07-DEC-1992 US 07/987130,07-DEC-1992 US 07/987130,07-
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C12N15/09,C12N5/10,C12N7/00,C12N9/22//(C12N5/10,C12R1:91), PC
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/mol_type="genomic RNA"
/db_xref="taxon:32630"
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Pred. No. 9.1e+05;
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PN JP 20034228
PD 12-DEC-2000
PP 11-MAY-2000
PP 01-MAY-1992 US
14-MAY-1992 US
15-OCT-1992 US
15-OCT-1992 US
                       Patent: JP : GERON CORP
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CC
FH Key
FT source
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PI JAME
PC C12N
PC A61K
PC A61K
PC A61K
PC A61F
PC A61F
                                          1 (bases 1 to 20)
Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 41 09-OCT-2001;
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BD071075.1 GI:22616678
JP 2001517929-A/41.
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Modulation of mammalian
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1 (bases 1 to 20)
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(C12N15/09,C12R1:93),C12N15/00,C12N5/00,A61K37/48,(C12N15/00,
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11-MAY-1992 US 07/88268),14-MAY-1992 US MAY-1992 US 07/88273,14-MAY-1992 US 07/88286,14-MAY-1992 US 07/88286,14-MAY-1992 US 07/882886,14-MAY-1992 US 07/882889,14-MAY-1992 US 07/882889,14-MAY-1992 US 07/882889,14-MAY-1992 US 07/882922,14-MAY-1992 US 07/882922,14-MAY-1992 US 07/882922,14-MAY-1992 US 07/882921,14-MAY-1992 US 07/882921,14-MAY-1992 US 07/88292,14-MAY-1992 US 07/88292,00-DEC-1992 US 07/96302,00-DEC-1992 US 07/96302,00-DEC-1992 US 07/963122,00-DEC-1992 US 07/963122 US 07/963122
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C12N15/09,C12N5/10,C12N7/00//A61K38/43,A61K39/125,A61K39/13,
A61K39/135,
A61K39/145,A61K39/21,A61K39/23,A61K39/245,A61K39/29,A61K48/00,
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/mol_type="genomic RNA"
/db_xref="taxon:32630"
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07/882823 PR

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07/8844073 PR

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TITLE
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                                                                                                                                                               Shay, J.W., Wright, W.E., Piatyszek, M.A., Modulation of mammalian telomerase by patent: JP 2001517929-A 46 09-OCT-2001;
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BD071080.1 GI:2261
JP 2001517929-A/46.
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C07K14/00,A61K38/16,C12Q1/68
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09-APR-1996 US 08/630019
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09-OCT-2001
                                                      /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                          Location/Qualifiers
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mol_type="genomic DNA"
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/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC)
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Location/Qualifiers
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9.1e+05;
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RESULT 205
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JF 2001321190-A/1423.
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A method of arraying genome
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                                                                        A method of arraying genome clone
Patent: JP 2001321190-A 1423 20-NOV-2001;
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                                                              INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH, YUGENKAISHA
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Conservative
     Artificial Sequence
JP 2001321190-A/1423
20-NOV-2001
12-MAR-2001 JP 2001068285
                                                                                                                                                                                                                                                                                            2.0%; Solitarity 100.0%; I Conservative 0;
EIICHI SOEDA
                                                                                                                                                                                                                                                                                                                                                                                                                                      Description of Artificial Sequence: Synthetic DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Artificial Sequence JP 2001321190-A/687
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2-MAR-2001 JP 2001068285
                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                          /organism='Artificial
                                                                                                                                                                                                                                                                                                                                                 /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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ches 0;
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9.1e+05;
hes 0;
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ORGANISM
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GENESENSE TECHNOLOGIES INC
OS Homo sapiens (human)
PN JP 2002501743-A/14
PD 22-JAN-2002
PF 29-JAN-1999 JP 20005294
PF 29-JAN-1998 US 60/07
PI JIM A WRIGHT,AIPING H Y
PC C12N15/09,A61K31/711,A6
PC (A61K31/711,A61K45:00),
442 TGGAATACTTTT 453
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PI JIM A WRIGHT, AIPING + YOUNG, YOON S LEE
PC C12N15/09, A61K31/711, A61K48/00, A61P35/00, A61P35/04, C07H21/04//
PC (A61K31/711, A61K45:00), (A61K48/00, A61K45:00), C12N15/00 CC
Cligonucleotide sequence complementary to thioredoxin gene or CC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wright, J.A., Young, A.H. and Lee, Y.S.
Oligonucleotide sequence complementary to thioredoxin gene or thioredoxin reductase gene and utilization thereof for controlling Patent: JP 2002501743-A 14 22-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (human)
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C12N15/00,
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JP 2002501743-A/14.
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Description of Artificial Sequence:Synthetic DNA
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30-JAN-1998 US 60/0731
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                                                                                                                                                                                                                                                                                /organism="Homo sapiens'
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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100.0%; Pr
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9.1e+05;
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ACCESSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     source
artificial sequences.

21 (bases 1 to 20)

22 Nagamine,K., Hase,T. and Notomi,T.

Method of synthesizing single-stranded nucleic acid

Patent: WO 0234907-A 22 02-MAY-2002;

EIKEN CHEMICAL CO LTD,KENTARO NAGAMINE,TETSU HASE,TSUGUNORI NOTOMI

OS Artificial Sequence

WO 0234907-A/22

PD 02-MAY-2002

PF 26-CCT-2001 WO 2001JP009452

PF 26-CCT-2001 WO 2001JP009452

PR 27-CCT-2000 JP 00P 328219

PR XENTARO NAGAMINE,TETSU HASE,TSUGUNORI NOTOMI

PC C12N15/10,C120/68

CC Description of Artificial Sequence:Outer primer(Forward) FH

LOCATION Qualifiers

FT source /organism='Artificial Sequence'.
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Method of
BD170190
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Methods for analyzing animal

BD135497

BD135497.1 GT:23230442

JP 2002504814-A/35.
                                                                                                                                                                                                                    BD170190.1 GI:27876002
WO 0234907-A/22.
synthetic construct
synthetic construct
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S, RICHARD WALES,
GRAHAM STUART PLASTOW
                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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30-MAY-1997 GB 9711214.8,31-JAN-1998 GB
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                                                                                                                                                                                                                                                                                  90 20 bp DNA line of synthesizing single-stranded nucleic
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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0; Mismatches
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9.1e+05;
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12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (04-AUG-2001) Akira Horii, Tohoku University Schu Medicine, Molecular Pathology; 2-1 Sziryomachi, Aoba-ku, S. Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp, Tel:81-22-717-8042, Fax:81-22-717-8047)
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Genomics 74 (1), 55-70 (2001)
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                                                                         AB069312
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A BAC-based STS-content map spanning a 35-Mb region of human
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sts-R90NSR obtained from clones B133N1, B369A24, B341F17,
B262P12, B341F17, B334L24, B128O2, B90N5, B229F2, Human
BAC library RPCI-11"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 9.1e+05;
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9.1e+05;
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A BAC-based STS-content map spanning a 35-Mb region of human chromosome 1p35-p36
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                                                                                                                                                                                      Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 4590 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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sts-stSG13123 obtained from clones B157K8, B14F15, B:
B251F9, Human BAC library RPCI-11"
                                                                                                                   /organism="Homo sapiens"
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RESULT 212

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11; Conserv
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Sequence 9848 from Patent
AX632806
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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BD242529/c
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AR058692
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             CELLOMICS INC
OS Artificia
PN JP 200252
PD 03-SEP-20
PF 29-OCT-19
PR 30-OCT-19
                                                                                 artificial sequences.

1 (bases 1 to 12)

Guiliano, K.A., Bright, G., Olson, K. and Tencza, S. A system for cell based screening
Patent: JP 2002528136-A 35 03-SEP-2002;
                                                                                                                                                                          BD242529
BD242529.1 GI:33052299
JP 2002528136-A/35.
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AR058692
                                                                                                                                                  synthetic construct synthetic construct
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Chee, M., Cronin, M.T., Fodor, S.P.A., Huang, X.X., Hubbell, E.A.,
Lipshutz, R.J., Lobban, P.E., Morris, M.S. and Sheldon, E.L.
Arrays of nucleic acid probes on biological chips
Patent: US 5837832-A 269 17-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Н
                                                                                                                                                                                                             A system for cell based screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Method for determining homeostasis of the skin
Patent: WO 02053774-A 9848 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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                                    Artificial Sequence
JP 2002528136-A/35
03-SEP-2002
29-OCT-1999 JP 2000579780
30-OCT-1998 US 60/106308,26-MAY-1999 US
3TH A GUILIANO,GARY BRIGHT,KEITH OLSON,SARAH
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 3.3e+06;
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AX766780/c
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 organism="synthetic construct"/
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Giuliano, K.A. and Kapur, R.
A system for cell-based screening
Patent: EP 1314980-A 69 28-MAY-2003;
Cellomics, Inc. (US)
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Sequence 69 from Patent EP1314980.
AX766780
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Giuliano,K. and Kapur,R.
Gystem for cell-based screening
Patent: US 6416959-A 69 09-JUL-2002;
Location/Qualifiers
                                                                                             synthetic construct artificial sequences.
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PC C12N15/09,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12Q1/02,C12Q1/
PC 37,G01N33/15,
PC G01N33/50,C12N15/00,C12N5/00
PC G01N33/50,C12N15/00,C12N5/00
PC G01N33/50,C12N15/00,C12N5/00
                                                                                                                             synthetic construct
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G01N33/50,C12N15/00,C12N5/00
Description of Artificial Sequence:
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Location/Qualifiers
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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artificial sequences.
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Direct Submission
Submitted (23-OCT-2003) Balzergue S., UMRGV, INRA/CNRS, 2 rue
Gaston Cremieux, 91057 Evry cedex, FRANCE
PCR was performed on DNA from transformants of Arabidopsis thaliana
                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                   AJ598336.1 GI:37947964
right border; T-DNA flanking sequence.
Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
                                                                                                                                                                                Brunaud, V., Balzergue, S., Dubreucq, B., Aubourg, S., Samson, F., Chauvin, S., Bechtold, N., Cruaud, C., DeRose, R., Pelletier, G., Lepiniec, L., Caboche, M. and Lecharny, A.
                                                                                                                                                                                                                 Brunaud, V.,
                                                                                                                                                                                                                                                                                                                                                                                                    AJ598336 13 bp DNA linear PLN 23-OCT-200
Arabidopsis thaliana T-DNA flanking sequence, right border, clone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent: WO 02070755-A 508 12-SEP-2002;
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BO Rep. 3 (12), 1152-1157 (2002)
                                                                                   (bases 1 to 13)
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/db_xref="taxon:32630"
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100.0%; Pred. No. 3.3e+06;
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; Pred. No. 3.3;
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. 3.3e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plants from INRA (Versailles). The DNA fragment(s) resulting from the PCR were directly sequenced from the left or the right border to determine the genomic sequence flanking the insertion. T-DNA derived sequences were removed. Information to order the corresponding mutant line and a link to a database providing a graphical display of the insertion site are available at http://dbsgap.versailles.inra.fr/publiclines/. This sequence has been generated in the framework of the French plant genomics program 'Genoplante' (http://www.genoplante.com and http://genoplante-info.infobiogen.fr).
A88769
Sequence 917 from Patent
A88769
A88769.1 GI:6737339
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Sequence 97 from Patent WO9502051.
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                                                                                                                                                                                                                        l Similarity
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BEYSEN, W.

A PHARMACEUTICAL COMPOSITION COMPRISING ANTISENSE-NUCLEIC ACID FOR PREVENTION AND/OR TREATMENT OF NEURONAL INJURY, DEGENERATION AND CELL DEATH AND FOR THE TREATMENT OF NEOFLASMS PATENT: WO 9502051-A 97 19-JAN-1995; BICKNOSTIK GES FUER BIOMOLEKUL (DE)
Other publication AU 7345694 950206.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unclassified.
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/mol_type="genomic_DNA"

/culTivar="Wassillewskija"

/db_xref="taxon:3702"

/clone="466D02"
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                                                                                                                                                                                                                                                                                                                /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon;32644"
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he right border
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                                    1 (bases 1 to 14)
Regan, J.W., Gil, D.W. and Woodward, D.F.
Nucleic acid encoding a human EP prostaglandin receptor
Patent: US 6395878-A 1 28-MAY-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                          l (bases 1 to 14)
Regan, J.W., Gil, D.W. and Woodward, D.F.
Nucleic acid encoding a novel human EP
Patent: US 5716835-A 1 10-FEB-1998;
Location/Qualifiers
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11; Conserv
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Brysch, W. and Schlingensiepen, K.

AN ANTISENSE OLIGONICLEOTIDE PREPARATION METHOD

Patent: WO 9833904-A 917 06-AUG-1998;

BIOGNOSTIK GES (DE), BRYSCH WOLFGANG (DE)
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AR372103
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       /organism="unknown"
/mol_type="genomic
                                                                                                                                                                                                                                                                                                                                             /organism="unknown"
/mol_type="unassione
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/db_xref="taxon:32644"
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                                                                                                               Schlingensiepen, K.H. and Brysch, W.

An antisense oligonucleotide preparation method

L Patent: JP 2001511000-A 917 07-AUG-2001;

BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH

OS Unknown

JP 2001511000-A/917

PD 07-AUG-2001

PF 30-JUR-1998 JP 1998532533

PF 31-JAN-1997 EP 97101531.8

PR 31-JAN-1997 EP 97101531.8

PR 31-JAN-1997 EP 97101531.9

PC C12N15/11,007H21/04,A61K31/70

CC An antisense oligonucleotide preparation method FH Ke
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BD066282
BD066282.1 GI:22611885
JP 2001511000-A/917:
unidentified
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Use of csf-1 inhibitors
Patent: WO 0130381-A 8 (AT)
Hofbauer, Reinhold (AT)
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Sequence 8 from Patent WO0130381.
AX133710
AX133710.1 GI:14139720
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                                                                                              30-JAN-1998 JP 1998532533
31-JAN-1997 EP 97101531.8
KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
C12N15/11,C07H21/04,A61K31/70
An antisense oligonucleotide preparation method
Location/Qualifiers
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                    /organism='Unknown'
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/db_xref="taxon:32630"
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Pred. No. 3.3e+06;
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BD201796/c
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V Enzymatic nucleic acid treatment of di
to hepatitis C virus infection.
BDZ09325.
BDZ09325.1 GI:33019095
BDZ09325.191-A/2915.
Unidentified
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PC A61P35
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BD201796.1 GI:33011566
JP 2002509721-A/4822.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                               concerning molecule
participating in vasculogenic response
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27-MAR-1998 US 60/079678
PAMELA A PAVCO, ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT,
                                                                                                                                                                                                                                                                                                                                                                                      source
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                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                    /organism='Homo sapiens
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0; Mismatches
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OS Hepatitis virus (hepatitis C virus)
PN UP 2002512791-A/2915
PD 08-MAY-2002
PF 26-APR-1999 JP 2000545991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l (bases 1 to 14)
Blatt, L., Mcswiggen, J.A., Roberts, E., Pavco, P.A. and Macejak, D.
Enzymatic nucleic acid treatment of diseases or conditions rela
to hepatitis C virus infection
Patent: JP 2002512791 A 2915 08-MAY-2002;
                                                                                                                                                                                                                                                                                            Sequence
AR033266
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PI II
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                                                                                                                                                                      Draper, K.G. Method and reagent for inhibiting hepatitis C virus replication Patent: US 5869253-A 32 09-FEB-1999;
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27-APR-1998 US 60/083217,18-SEP-1998 US 60/100842 PR

25-FEB-1999 US 09/257608,23-MAR-1999 US 09/274553 PI

LAWRENCE BLATT,JAMES A MCSWIGGEN,ELISABETH ROBERTS,PAMELA A PI
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                                                                                                                                                                                                                                                       Unknown.
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                       AGGCTGAGCCC 369
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A61K37/66,
C12N15/00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="unidentified"
/mol_type="genomic RNA"
/db_xref="taxon:32644"
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                                                                                                                 /organism="unknown"
/mol_type="unassigned DNA"
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Pred. No.
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RESULT 230 AR056295

PAT 18-FEB-2000

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AR056532
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AR056484
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Sequence
AR056484
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Sequence
AR056295
Draper, K.G.
Intercellular adhesion molecule-1 (ICAM-1)
Patent: US 5837542-A 736 17-NOV-1998;
                                                                                                     Sequence 736 from patent AR056532 AR056532.1 GI:5982109
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                                  1 (bases 1 to 15)
Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
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Intercellular adhesion molecule-1 (ICAM-1) ribozymes
Patent: US 5837542-A 499 17-NOV-1998;
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Grimm, S., Stinchcomb, D.T., McSwiggen, J.,
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Grimm, S., Stinchcomb, D.T., McSwiggen, J.,
                                                                                                                                      AR056532
                                                                                                                                                                                                                                                                                                                                                  Patent: US 5837542-A 688 17-NOV-1998;
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                                                         Unclassified.
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Similarity 100.0%; Pred. No. 3.3e+06;
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                                                                                                                                                                                                                                                                                                             organism="unknown"
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                                   Sullivan, S.
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Haselkorn, R. and Gornicki, P.
Acetyl-Coa, Carboxylase compositions and
Patent: US 5910626-A 31 08-JUN-1999;
                                                                                                                                                          Method and reagent for inhibiting hepatitis C virus replication Patent: US 6132966-A 32 17-OCT-2000;
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Draper, K.G.
                                                                                                                                                                                                                                Unknown
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                                                                                                                                                                                                                                                                                      32 from patent US
                                                                                                                        /organism="unknown"
/mol_type="unassigned
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/mol_type="unassigned DNA"
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AR114053
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AR114290
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 15)
Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
Draper,K.G.
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                                                                                                                                                                                               11;
                                                                 AR114290
Sequence 736 from patent
AR114290
                                                                                                                                                                                                                                                                                        Ribozyme treatment of diseases or conditions related to levels intercellular adhesion molecule-1 (ICAM-1) Patent: US 6132967-A 688 17-OCT-2000; Location/Qualifiers
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                    Unknown
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S. and
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          Unclassified
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(bases 1 to 15)
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688 from patent US 6
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Pred. No. 3.3e+06;
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AR132197
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1 (bases 1 to 15)

Stinchcomb,D.T., Jarvis,T. and McSwiggen,J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 622 27-FEB-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ribozyme treatment of diseases or conditions related to levels of intercellular adhesion molecule-1 (ICAM-1)
Patent: US 6132967-A 736 17-OCT-2000;
Location/Qualifiers
                                                                                                     Unclassified.

1 (bases 1 to 15)
Stinchcomb,D.T., Jarvis,T. and McSwiggen,J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 623 27-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                 l Similarity
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h 1.8%; Score 11; DB 6; L
Similarity 100.0%; Pred. No. 3.3e+06;
11; Conservative 0; Mismatches 0;
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nilarity 100.0%;
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                                                          /organism="unknown"
/mol_type="unassigned DNA"
                                                                                             Location/Qualifiers
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AR133396
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ORGANISM
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AR132200
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AR133396
AR133396.1
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Sequence
AR132199
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Stinchcomb, D.T., Jarvis, T. and McSwiggen, J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 625 27-FEB-2001;
Location/Qualifiers
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AR132200
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1 (bases 1 to 15)
Stinchcomb,D.T., Jarvis,T. and McSwiggen,J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 624 27-FEB-2001;
Location/Qualifiers
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Unclassified
                    Unknown.
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625 from patent US 6194150.
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/mol_type="unassigned DNA"
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                                          GI:14122301
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AR133398
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Best Local (
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                                                                      Unclassified.

1 (bases 1 to 15)

1 (bases 1 to 15)

Stinchcomb, D.T., Jarvis, T. and McSwiggen, J.

Nucleic acid based inhibition of CD40

Patent: US 6194150-A 1823 27-FBB-2001;

Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                             Unclassified.

1 (bases 1 to 15)
Stinchcomb, T., Jarvis, T. and McSwiggen, J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 1822 27-FEB-2001;
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AR133397
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Stinchcomb, D.T., Jarvis, T. and McSwiggen, J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 1821 27-FEB-2001;
Location/Qualifiers
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nilarity 100.0%;
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                                                      /organism="unknown"
/mol_type="unassigned
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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41 AATTCAAAAAT 51

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I39403/c
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139402/c
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1 (bases 1 to 15)
Pogo,A.O. and Chaudhuri,A.
DNA encoding Duffy 9pd protein
Patent: US 5578714-A 4 26-NOV-1996;
Location/Qualifiers
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I30019
                                                          I39403
Sequence 441 from patent
I39403
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Sullivan,S., Draper,K.G., McSwiggen,J.
IL-5 targeted ribozymes
Patent: US 5616488-A 440 01-APR-1997;
Location/Qualifiers
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Sequence 440 from patent US 5616488.
139402
139402.1 GI:2083882
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(bases 1 to 15)
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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157495
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Sullivan,S., Draper,K.G., McSwiggen,J. and Stinchcomb,D.T.
IL-5 targeted ribozymes
Patent: US 5616488-A 442 01-APR-1997;
Location/Qualifiers
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Sequence 442 from patent US 5616488.
139404
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IL-5 targeted ribozymes
Patent: US 5616488-A 441 01-APR-1997;
Location/Qualifiers
                                                                                                                  Draper, K.G.
Enzymatic RNA molecule targeted against Hepatitis C virus
Patent: US 5610054-A 32 11-MAR-1997;
                                                                                                                                                                      Unknown
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                       Conservative
                                                                                                                                                                                                                          32 from patent US
                                                                      /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Aharinejad,S.
Use of csf-1 inhibitors
Patent: WO 0130381-A 11 03-MAY-2001;
Hofbauer, Reinhold (AT)
Location/Qualifiers
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171878
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unidentified
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synthetic construct
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                                          AX633447.1 GI:28469061
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ilarity 100.0%; Pred. No. 3.3e+06;
Conservative 0; Mismatches 0;
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4 from patent US 5683696.
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/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                              /note="Primer"
                                                                                                                                                                                                                                                                                         organism="synthetic construct"
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Pred. No. 3.3e+06;
                                                                    EP1260586.
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unidentified unidentified unclassified.
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AX633547
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RIBOZYME PHARMACEUTICALS, INC. (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stinchcomb, D.T., Dudycz, L.W., Chowrira, B., Grimm, S., Direnzo, A., Karpeisky, A., Draper, K.G., Kisich, K., Matulic-Adamic, J., Mcswiggen, J.A., Modak, A., Pavco, P., Beigelman, L., Sullivan, S.M., Sweedler, D., Thompson, J.D., Tracz, D., Usman, N., Wincott, F.E. and
                                                      AX633547.1
                                                                                             AX633547
                                                                                                                                                                                                                                                                                                                                             Patent: EP 1260586-A 627 27-NOV-2002; RIBOZYME PHARMACEUTICALS, INC. (US) Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                              Stinchcomb,D.T., Dudycz,L.W., Chowrira,B., Grimm,S., Direnzo,A., Karpeisky,A., Draper,K.G., Kisich,K., Matulic-Adamic,J., Mcswiggen,J.A., Mcdak,A., Pavco,P., Beigelman,L., Sullivan,S.M., Sweedler,D., Thompson,J.D., Tracz,D., Usman,N., Wincott,F.E. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 627 from Patent AX633488
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                                                                                                                                                                                                                                                                                        /organism="unidentified"
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/db_xref="taxon:32644"
                                                        GI:28469161
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Pred. No. 3.3e+06;
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Patent: EP 1260586-A 686 27-NOV-2002;
RIBOZYME PHARMACEUTICALS, INC. (US)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Method and reagent for inhibiting the expression of disease related
                                                                                                     Sequence 2828 from Patent AX635689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stinchcomb, D.T., Dudycz, L.W., Chowrira, B., Grimm, S., Direnzo, A., Karpeisky, A., Draper, K.G., Kisich, K., Matulic-Adamic, J., Mcswiggen, J.A., Modak, A., Pavco, P., Beigelman, L., Sullivan, S.M., Sweedler, D., Thompson, J.D., Tracz, D., Usman, N., Wincott, F.E. and
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                 unidentified unclassified
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RIBOZYME PHARMACEUTICALS, INC. (US)
Location/Qualifiers
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/mol_type="unassigned RNA"
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/mol_type="unassigned RNA"
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Sequence 2830 from Patent
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AX635691.1 GI:28471305
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unclassified

1 (bases 1 *
                                                                 BD206999
BD206999.1 GI:33016769
JP 2002512791-A/589.
                                                                                                         Enzymatic nucleic acid treatment of diseases to hepatitis C virus infection.
BD206999
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11; Conserv
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/mol_type="unassigned RNA"
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/mol_type="unassigned RNA"
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RESIDENT, Moswiggen, J.A., Roberts, E., Pavco, P.A. and Macejak, D. Blatt, L., Moswiggen, J.A., Roberts, E., Pavco, P.A. and Macejak, D. Blatt, L., Moswiggen, J.A., Roberts, E., Pavco, P.A. and Macejak, D. Blatt, L., Moswiggen, J.A., Roberts or conditions related to hepatitis C virus infection

AL Patent: JP 2002512791-A 1943 08-MAY-2002;

RIBOZYME PHARMACEUTICALS INC

OS Hepatitis virus (hepatitis C virus)

PN JP 2002512791-A/1943

PN JP 2002512791-A/1943

PN JP 2002512791-A/1943

PF 26-APR-1999 UP 200545991

PF 26-APR-1999 UP 60/083217, 18-SEP-1998 US 60/100842 PR 27-APR-1998 US 60/083217, 18-SEP-1998 US 60/100842 PR 27-APR-1998 US 60/083217, 18-SEP-1998 US 09/274553 PI LAWRENCE BLATT, JAMES A MCSWIGGEN, ELISABETH ROBERTS, PAWELA A PI PAVCO,

PI DENNIS MACEJAK

PC C11N15/00

CC Enzymatic nucleic acid treatment of diseases or conditions CC Enzymatic nucleic acid treatment of diseases or conditions CC Enzymatic nucleic acid treatment of diseases or conditions CC
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Enzymatic nucleic acid treatment of diseases or conditions related
to hepatitis C virus infection
Patent: JP 2002512791-A 589 08-MAY-2002;
RIBOZYME PHARMACEUTICALS INC
OS Hepatitis virus (hepatitis C virus)
PD JP 2002512791-A/589
PD 08-MAY-2002
PD 08-MAY-2002
PD 08-MAY-2002
PP 26-APR-1999 US 60/003217,18-SEP-1998 US 60/100842 PR
27-APR-1998 US 60/003217,18-SEP-1998 US 09/274553 PI
LAWRENCE BLATT, JAMES A MCSWIGGEN, ELISABETH ROBERTS, PAMELA A PI
PAVCO,
PI DENNIS MACEJAK
PC C12N9/00,A61K31/7105,A61K38/21,A61K48/00,A61P31/12,C12N15/09,
PC C12N15/06
CC Enzymatic nucleic acid treatment of diseases or conditions CC
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JP 2002512791-A/1943.
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hepatitis C virus infection.
Key Location/Out
                                            Enzymatic nucleic acid treatment related to
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/mol_type="genomic RNA"
/db_xref="taxon:32644"
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                                                                                                                                                      Sequence 4
AR089188
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A66853
                                          Homogeneous methods for nucleic acid
Patent: US 5994056-A 4 30-NOV-1999;
Location/Qualifiers
                                                                           1 (bases 1 to 16) Higuchi, R.G.
                                                                                                            Unknown
                                                                                                                                                                                                                                                                                                                                                                                          Stuyver, L., Rossau, R. and Maertens, G. METHOD FOR TYPING AND DETECTING HBV Patent: WO 9740193-A 20 30-OCT-1997; INNOGENETICS NV (BE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          unidentified unidentified
                                                                                                                         Unknown
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20 from Patent WO9740193.
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/mol_type="genomic RNA"
/db_xref="taxon:32644"
         /organism="unknown"
/mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                   /mol_type="unassigned DNA"
/db_xref="taxon:32644"
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1.8%; Score 11; DB 6; L ilarity 100.0%; Pred. No. 3.3e+06; Conservative 0; Mismatches 0;

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Chakraborty, P.R., Dashkevicz, M., Elbrecht, A., Feighner, S.D., Liberator, P.A. and Profous-Juchelka, H.
Eimeria praecox 16S rDNA probes
Patent: US 5449768-A 31 12-SEP-1995;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                      Sequence 31 from patent US I14457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unclassified.
1 (bases I to 16)
Higuchi, R.G.
Methods and devices for hemogeneous nucleic acid amplification and
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                     Sequence 31 from patent US I27300
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          GI:1818076
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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100.0%; Prr
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100.0%; Pred. No. 3.3
tive 0; Mismatches
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Chakraborty, P.R., Dashkevicz, M., Elbre,
Liberator, P.A. and Profous-Juchelka, H.
Eimeria tenella 16s rNNA probes
Patent: US 5563256-A 31 08-OCT-1996;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 145 from patent 127973 127973.1 GI:1818749
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Apple, R.J., Erlich, H.A., Griffith, R.L. and Schamptle, R.J., Erlich, H.A., Griffith, R.L. and Schamethods and reagents for HLA DRbeta DNA typing Patent: US 5567809-A 145 22-OCT-1996;
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                                                                    1 (bases 1 to 16)
Apple, R.J., Erlich, H.A., Griffith, R.L. a
Apple, R.J., Erlich, H.A., Griffith, R.L. a
Methods and reagents for HLA DRbeta DNA
Patent: US 5567809-A 152 22-OCT-1996;
Location/Qualifiers
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                                   /organism="unknown"
/mol_type="unassigned
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JS 5567809.
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3.3e+06;
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PD 11-C
PF 28-C
PR 02-C
PI TAKC
PC C12:
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127982
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1 (Dases 1 to 16)

Shibata,T., Noguchi,Y. and Ymashita,M.

A gene coading a cyclic lopopeptide acylase and an expression

Patent: WO 0102585-A 42 11-JAN-2001;

FUJISAWA PHARMACEUTICAL CO LTD, TAKASHI SHIBATA, YUJI NOGUCHI, MICHIO
                                                                                                                                                                                                                                                                                                                                                     BD093179
BD093179.1 GI:2
WO 0102585-A/42.
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Apple.R.J., Erlich,H.A., Griffith,R.L. and Scharf,S. Methods and reagents for HLA DRbeta DNA typing Patent: US 5567809-A 154 22-OCT-1996;
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1.8%; Score 11; DB 6; Le ilarity 100.0%; Pred. No. 3.3e+06; Conservative 0; Mismatches 0;
                                                                                                                         28-JUN-2000 WO 2000JP004285
02-JUL-1999 JP 99P 189644
TAKASHI SHIBATA, YUJI NOGUCHI, MICHIO YMASHITA
C12N15/55,C12N1/21,C12N9/14
Oligonuclectide designed to act as sequencing
Location/Qualifiers.
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WO 0102585-A/42
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coading a cyclic lopopeptide
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154 from patent US 5567809.
                                                              /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mol_type="unassigned DNA"
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                                    RESULT 271
AR039559
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ORGANISM
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AR039559
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AR039557
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A42338
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Stinchcomb, D.T. and McSwiggen, J.A.
Interleukin-2 receptor gamma-chain ribozymes
Patent: US 5807743-A 405 15-SED-1998;
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Gusterson, B.A., Crompton, M.R., Mitchell, P.J.,
Kamalati, T., Page, M.J. and Spence, P.
PROTEIN TYROSINE KINASE AND LIGANDS THEREOF
Patent: WO 9502057-A 10 19-JAN-1995,
CANCER RES INST (GB)
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                /organism="unknown"
/mol_type="unassigned DNA'
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from Patent WO9502057
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          17 bp I
from patent US 5807743.
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AR057431
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AR039561
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Best Local Similarity
Matches 11; Conserv
                                                                           TITLE
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1 (bases 1 to 17)

Stinchcomb,D.T. and McSwiggen,J.A.

Interleukin-2 receptor gamma-chain ribozymes

Patent: US 5807743-A 407 15-SEP-1998;
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AR039561
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Stinchcomb.D.T. and McSwiggen,J.A.
Interleukin-2 receptor gamma-chain ribozymes
Patent: US 5807743-A 409 15-SEP-1998;
Location/Qualifiers
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AR057431
AR057431.1 GI:5983008
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                                                           Draper, K.G.
Intercellular adhesion molecule-1 (ICAM-1) ribozymes
Patent: US 5837542-A 1635 17-NOV-1998;
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                                                                                                                                         Unknown
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409 from patent US 5807743.
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
          /organism="unknown"
/mol_type="unassigned DNA"
                                                   Location/Qualifiers
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100.0%; Pred. No.
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RESULT 275
AR057566
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AR057487
LOCUS
RESULT 276
AR057626
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Intercellular adhesion molecule-1 (ICAM-1) ribozymes
Patent: US 5837542-A 1691 17-NOV-1998;
Location/Qualifiers
1. .17
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
Draper,K.G.
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                                                                                                                                                           Sequence 1770 from patent AR057566
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                                                                                                                                                                                                              1 (bases 1 to 17)
Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
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                                                                                                                                                                                                                                    Unclassified.
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                                                                                                                                    /organism="unknown"
/mol_type="unassigned
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                                                                                  Score 11; DB 6; L; Pred. No. 3.3e+06; 0; Mismatches 0;
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S. and
Draper,K.G.
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AR057632
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AR057626
AR057626.1 GI:5983203
                                          1 (bases 1 to 17) Grimm, S., Stinchcomb, D.T., McSwiggen, J., Sullivan, S.
                                                                                                                           Sequence 1891 from patent AR057687
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                                Draper, K.G.
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Patent: US 5837542-A 1830 17-NOV-1998;
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Unknown
                                                                                                                AR057687.1 GI:5983264
                                                                                                                                                   AR057687
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                                                                   Inclassified
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Patent: US 5837542-A 1836 17-NOV-1998;
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/mol_type="unassigned DNA"
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Pred. No. 3.3e+06;
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Best Local Similarity 100.0%; Pred. No. 3.3e+06;
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                                                                                                                                                           Patent: US 5837542-A 1968 17-NOV-1998;
                                                                                                                                                                                 Draper, K.G.
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Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
                                                                                                                                                                                                                              Unknown
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Patent: US 5837542-A 1894 17-NOV-1998;
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Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
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                                                                  Similarity
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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AR057782
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Sequence 1984 from part
AR057780
AR057780.1 GI:5983357
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Unclassified.

1 (bases 1 to 17)
Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
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Grimm, S., Stinchcomb, D.T., McSwiggen, J.,
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Patent: US 5837542-A 1981 17-NOV-1998;
    Location/Qualifiers
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Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
                                  Unknown
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/mol_type="unassigned DNA"
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Intercellular adhesion molecule-1 (ICAM-1) ribozymes
Patent: US 5837542-A 1986 17-NOV-1998;
Location/Qualifiers
                                                                                                1 (bases 1 to 17)
Grimm, S., Stinchcomb, D.T., McSwiggen, J.,
Draper, K.G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribozyme treatment of diseases or conditions related to levels of intercellular adhesion molecule-1 (ICAM-1)
Patent: US 6132967-A 1635 17-OCT-2000;
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                                               Ribozyme treatment of diseases or conditions related to levels of intercellular adhesion molecule-1 (ICAM-1) Patent: US 613957-A 1691 17-OCT-2000; Location/Qualifiers
                                                                                                                                                      Unknown
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
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1635 from patent US
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AR115390
Sequence 1836 from
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
Draper,K.G.
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Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
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                                                                                                                        1.8%; Score 11; DB 6; Le clarity 100.0%; Pred. No. 3.3e+06; Conservative 0; Mismatches 0;
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larity 100.0%; Pred. No.
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
Draper,K.G.
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Ribozyme treatment of diseases or conditions related to levels of intercellular adhesion molecule-1 (ICAM-1)
                              1 (bases 1 to 17) Grimm, S., Stinchcomb, D.T., McSwiggen, J., Sullivan, S.
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Draper,K.G.
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                                                                                                            Ribozyme treatment of diseases or conditions intercellular adhesion molecule-1 (ICAM-1) Patent: US 6132967-A 1981 17-OCT-2000;
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
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(larity 100.0%; Pred. No. 3.3e+06;
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/mol_type="unassigned DNA"
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AR142556.1 GI:15103842
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Ribozyme treatment of diseases or conditions related intercellular adhesion molecule-1 (ICAM-1) Patent: US 6132967-A 1984 17-OCT-2000;
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Grimm, S., Stinchcomb, D.T., McSwiggen, J., Sullivan, S.
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Bribozyme treatment of diseases or conditions intercellular adhesion molecule-1 (ICAM-1)

Patent: US 6132967, A 1986 17-OCT-2000;
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Grimm, S., Stinchcomb, D.T., McSwiggen, J., Sullivan, S. and
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Schaap, T. Cornelis., Kuiper, C.Maria. and
Coccidiosis polypeptide and vaccines
Patent: US 6203801-A 19 20-MAR-2001;
Location/Qualifiers
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Schasp, T.Cornelis, Kuiper, C.Maria. and Vermeulen, A.Nicolaas.
Coccidiosis polypeptide and vaccines
Patent: US 6203801-A 17 20-MAR-2001;
Location/Qualifiers
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Schaap, T.Cornelis, Kuiper, C.Maria. (
Schaap, T.Cornelis, Kuiper, C.Maria. (
Coccidiosis polypeptide and vaccines
Patent: US 6203801-A 15 20-MAR-2001;
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/mol_type="unassigned
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ACCESSION
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FEATURES
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AUTHORS
TITLE
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BD254479
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AR142562
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ACCESSION
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Best Local
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Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.

Regulation of repressor genes using nucleic a

Patent: JP 2002541795-A 2272 10-DEC-2002;

RIBOZYME PHARMACEUTICALS INC
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Sequence
AR142562
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Schaap, T. Cornelis., Kuiper, C. Maria. and
Coccidiosis polypeptide and vaccines
Patent: US 6203801-A 21 20-MAR-2001;
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11; Conserv
                                                                                                                                 C12P21/02,C12P21/02//A61K31/711,(C12N5/10,C12R1:91),(C12P21/02,
                                                                                                                                                                                                                                                                                                                                                   unclassified.
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JP 2002541795-A/2272.
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N JP 2002541795-A/2272

D 10-DEC-2002

F 11-APR-2000 JP 2000611654

R 12-APR-1999 US 60/129990

I LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN 12015/99, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC
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Location/Qualifiers
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/organism="unidentified|

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DEFINITION
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AUTHORS
TITLE
JOURNAL
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Search completed: March
Job time : 2790 secs
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Best Local S
Matches 11
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PN JP 2002541795-A/2273
PN JP 2002541795-A/2273
PD 10-DEC-2002
PF 11-APR-2000 JP 2000611654
PR 12-APR-1999 US 60/1129390
PR 12-APR-ENGE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN PC C12N15/09, A61X38/00, A61X48/00, A61P43/00, A61P43/00, C12N5/10, PC C12P21/02,
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Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.

Regulation of repressor genes using nucleic acid molecules

Patent: JP 2002541795-A 2273 10-DEC-2002;

RIBOZZME PHARMACEUTICALS INC
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A61K37/02,
(C12N5/00,C12R1:91)
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Location/Qualifiers
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/db_xref="taxon:32644"
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100.0%; Pred. No. 3.3e+06;
ative 0; Mismatches 0;
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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Run
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Ygapop 60.0, Ygapext 60.0
Fgapop 6.0, Fgapext 7.0
Delop 6.0, Delext 7.0
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Query SUMMARIES

45	44	43	42	C 41	40	w	w	w	c 36	w	w	33	c 32	31	30	N	c 28	N	N	c 25	N	N	N	c 21	N	μ.	ш	- سر	16	<u>.</u> د	C 14	μ.	 	11	10	ø	00	7	on	и		ი ა		<u>.</u>	Result
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489135 1M0319H1	486007 1M0313E1	559 1M0179	400362 1M0166C1	87854 1M0147K2	348201 1M0084F1	31739 1M0059D2	07763 IM0010F0	03578 1M0003H0	27699 NACL-	22764 HDN02-	51019 D09C0	558127 BX556	55876	13258 AU013258	64551 2M017	ZB34391 2M01	Z799394 2M005	784061 2M002	661787 IM054	17087 IM044	595242 1M040	95016 1M040	82154 1M037	98063 IM033	78491 1M029	47414 1M024	188 1M023	18201 1M019	Z394192 1M015	Z361152 1M010	Z309643 1M001	Z307864 1M001	37608 JMT	F316655 HDC	F305339 CLD1-	X029847 Šinal	885697 oj34f01.	26060 HNC23-1	88895 ad36b06	5064 ou61g11.x	584986 E011826-	50755 D06B10	98220 5009-0-4	22 5009-0-3	cription

ALIGNMENTS

REFERENCE	MSIN	SOURCE	VERSION	ACCESSION		DEFINITION	LOCUS	BM397622	XXXXXXX F
Eukaryota, Alveolata, Ciliophora, Oligohymenophorea, Hymenostomatida, Tetrahymenina, Tetrahymena. 1 (bases 1 to 14)	Tetrahymena thermophila	EST. Tetrahymena thermophila	BM397622.1 GI:18197675	BM397622	Tetrahymena thermophila cDNA, mRNA sequence.	5009-0-35-C02.t.2 Chilcoat/Turkewitz cDNA (large fraction)	BM397622 14 bp mRNA linear EST 17-JAN-2002		

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RESULT 2
BM398220/c
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SOURCE
ORGANISM
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                                                                                                                                      source
                                                                                                                                                               Contact: Turkewitz Ap
Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 60637,
Tel: 773 702 4374
Fax: 773 702 3172
Email: apturkew@midway.uchicago.edu
Seq primer: T3.
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Molecular Genetics and Cell Biology
University of Chicago
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BST from Tetrahymena thermophila, strain CU428.1, growing cells Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                       Tetrahymena thermophila Eukaryota; Alveolata; Ciliophora; Oligohymenophorea; Hymenostomatida; Tetrahymenina; Tetrahymena. 1 (base 1 to 14)
Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk Frankel,J. and Klobutcher,L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14 bp mRNA linear EST 17 5009-0-42-D11.t.1 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                       EST from Tetrahymena thermophila, strain CU428.1, growing cells Unpublished (2002)
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BM398220.1 GI:18198273
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            EST.
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/clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+, Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/mol_type="mRNA"
                                                                                                                   organism="Tetrahymena"
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                                                                  db_xref="taxon:5911"
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Percent Similarity:
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                                        US-09-966-880A-8 (1-198) x CA850755
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CA850755
CA850755.1 GI:33387548
EST.
Glycine max (soybean)
Glycine max (soybean)
Glycine max (soybean)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Soybean Genomics and Improvement Laboratory (SGIL) US Department of Agriculture (USDA), ARS, PSI Bldg.006, Rm 118, 10300 Baltimore Ave., Beltsville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 301 504 5750 Fax: 301 504 5728
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Alkharouf, N.W., Khan, R. and Matthews, B.F.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   infected by the soybean cyst nematode
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SerPheSerLeu 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                alkharon@ba.ars.usda.gov.
                                                                                                                                                                                                                                                                                                                                           /mol type="mRNA"
/cultivar="Peking"
/db_xref="taxon:3847"
/clone="D06Bl0"
                                                                                                                                                                                                                                               /tissue_type="Roots"
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/dev_stage="Seedlings"
/clone_lib="cDNA_Peking library 2, 4 day SCN3"
/clone_lib="cDNA_Peking library 2, 4 day SCN3"
/note="Vector: pBlusscript SK-; cDNA_clones from mRNA
extracted from Peking roots 2 and 4 days past invasion."
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                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Glycine max"
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Glycine max
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RESULT 5
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Insert Length: 15 Std Error: 0.00
Plate: 2 row: K column: 24
Seg primer: SP6; CATACGATTTAGGTGACACTATAG.
                     AI075064 16 bp mRNA linear EST 27-AUG-1998 ou61g11.x1 NCI CGAP Br2 Homo sapiens cDNA clone IMAGE:1632356 3' similar to TR:Q24348 Q24348 FIBRILLARIN; mRNA sequence.
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H. and Radelof,U.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Manranthaceae; Beta.

1 (bases 1 to 15)
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/lab_host="EMDH10B"
/clone_lib="MPIZ-ADIS-024-inflorescence"
/clone_vector: pcMVSPORT6; Site 1: Sall; Site 2: Not1;
/note="Vector: pcMVSPORT6; Site 1: Sall; Site 2: Not1;
CDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-Not1, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prof. Christian Jung; Sequence submission manaç
RZPD/GABI-Primary database: http://gabi.rzpd.de<sup>m</sup>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung: Sequence submission managed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SP6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note:
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/db_xref="taxon:161934"
/clone="024-002-K24"
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 16)
                                                                                                                                                           16 bp mRNA linear EST 28-OCT-1: qd35b06.xl Soares fetal heart NbHH19W Homo sapiens cDNA clone IMAGE:1731539 3' Similar to SW:SN22 HUMAN P51531 POSSIBLE GLOBAL TRANSCRIPTION ACTIVATOR SNF2L2 ;, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-GAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 712 Std Error: 0.00
Seg primer: -40m13 fwd. E7 from Amersham
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                    Homo sapiens
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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1 (bases 1 to 16)
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/clone lib="NCI CGAP Br2"
/note="Vector: pT7T3D-Pac (Pharmacia) with a modified
/note="Vector: pT7T3D-Pac (Pharmacia) with a modified
polylinker; 1st strand cDNA was prepared from pooled bulk
breast tumor tissue, and was then primed with a Not I -
oligo(dT) primer. Double-stranded cDNA was ligated to Eco
RI adaptors (Pharmacia), digested with Not I and cloned
into the Not I and Eco RI sites of the modified pT7T3
vector. This library is the normalized version of
NCI CGAP Br1.1. Library was constructed by Bento Soares
and M. Fatima Bonaldo. "
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/tissue_type="breast"
/lab_host="DH10B"
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/mol_type="mRNA"
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This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 543 Std Error: 0.00
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Unpublished (199)
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Mational Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                     Identification and initial characterization of 5000 expressed sequenced tags (BSTs) each from adult human normal and osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001)
                                                                                                                                                                                                                                                                                     Kumar, S., Connor, J.R., Dodds, R.A., Halsey, W., Van Horn, M., Mao, J.,
Sathe, G., Mui, P., Agarwal, P., Badger, A.M., Lee, J.C., Gowen, M. and
                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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HNC23-1-E1.R HNC (Human Normal of
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Contact: Sanjay Kumar
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//lab host="DH10B (ampicillin resistant)"
//clome_llb="Soares fetal heart_NhHH19W"
//clome_llb="Soares fetal heart_ThHH19W"
//clome_llb="Soares fetal heart_ThHH19W"
//clome_llb="Soares fetal heart_ThH19W"
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:1731539"
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709 Swedeland Road, 1
Tel: 610-270-7245
Fax: 610-270-5598
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cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 691 Std Error: 0.00
Seg primer: -40m13 fwd. ET from Amersham
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Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 19)
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Seq primer: T7
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Contact: Robert Strausberg, Ph.D.
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National Cancer Institute, Cancer Genome Anat
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/mol_type="mRNA"
/db_xref="taxon:9606"
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/lab_host="E.coli DH10 Branch
/clone lib="HNC (Human Normal Cartilage)"
note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
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Percent Similarity:
Best Local Similarity:
Query Match:
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AUTHORS
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                                                                                                                                                                                                                                                                                      Direct Submission
Submitted (06-JAN-2003) Genoscope
BP 191 91006 EVRY cedex - FRANCE (
- Web : www.genoscope.cns.fr)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                        Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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19 bp mRNA linear HTC 07-JAN-2003 Single read from an extremity of a full-length cDNA clone made from Anopheles gambiae total adult females. 3-PRIME end of clone FKOAAA44CD05 of strain 6-9 of Anopheles gambiae (African malaria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mosquito).
BX029847
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                                                                                                                                                   /db_xref="taxon:7165"
/clone="FKOAAA44CD05"
/plasmid="pME18S-FL"
/note="end : 3-PRIME"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /Clone_lib="NCI_CGAP_Lu5"
//Clone_lib="NCI_CGAP_Lu5"
//note="organ: lung; Vector: pT773D-Pac (Pharmacia) with a modified polylinker; lst strand cDNA was prepared from neuroemdocrine lung carcinoid, and was then primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                    /organism="Anopheles gambiae"
/mol_type="mRNA"
/strain="6-9"
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/lab_host="DH10B"
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/clone="IMAGE:1500217"
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Best Local Similarity:
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                                                                             CP316655 19 bp mRNA linear EST 15-AUG-HD--06-A14.gl OsHDAC1-overexpressing transgenic rice plasmid cD library (HD) Oryza sativa cDNA clone HD--06-A14, mRNA sequence. CP316655 CF316655.1 GI:33688416
                       Oryza sativa
Oryza sativa
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J.
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,
Large-scale Sequencing Analysis of Rice
Unpublished (2003)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
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Location/Qualifiers
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoldeae; Oryzeae; Oryza.
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: 82 31 330 6193
: 82 31 321 6355
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/cultivar="wackdong"
/db_xref="taxon:4530"
/clone="CLD1:-01-H03"
/tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Vector: pCR4-TOPO; Site 1: EcoRI; Leaf was incubated at 4 C(360uM/m-2sec-ī) for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR."
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/lab_host="E.coli DH10B"
/clone_lib="Rice_cold_treated_leaf_plasmid_cDNA_library
(CLD1)"
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Kim, J.S., Jun, K.M., Cheong, P. J., Kim, M. J., L.,
Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                           CF337608 19 bp mRNA linear EST 18-AUG-2003
JMT--08-C02.b1 AtJMT-overexpressing transgenic rice plasmid cDNA
library (JMT) Oryza sativa cDNA clone JMT--08-C02, mRNA sequence.
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Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                       Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                      Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                   Contact: Nahm B.H.
                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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CF337608.1 GI:33823602
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Ehrhartoideae; Oryzeae; Oryza.
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: 82 31 330 6193
: 82 31 321 6355
bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.Location/Qualifiers
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/mol type="mRNA"
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/clone="HD--06-A14"
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AUTHORS
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                Plate: 0010 row: F column: 16
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                             University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                                                       plasmid inserts
Unpublished (2000)
Contact: Robert B.
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AZ307864.1 GI:10347281
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                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0010 row: F column: 16
                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                               quality sequence stop: 19.
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(note="Vector: pCR4-T0P0; Site_1: EcoRI; Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was pepared from Arabidopsis Jasmoinate Carboxyl methyltransferase overexpression line."
                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
                                     /db_xref="taxon:10090"
/clone="UUGC1M0010F16"
                                                                                                                     organism="Mus musculus"
                                                                                                                                                           ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
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/clone="JMT--08-C02"
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/cultivar="Nackdong"
'lab_host="E. Coli strain XL10-Gold, T1-resistant,
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/clone_lib="Mouse 10kb plasmid UUGC1M library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson

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SOURCE
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1M0016E23F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0016E23 F, genomic survey secuence
                                                                                                                                                                                                                                                                                                                                        Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                                                                                                                  Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0016 row: E column: 23
                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
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gh quality sequence stop: 19.
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/organism="Mus musculus"
/mol_type="genomic DNA"
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Stdn: 16
Plate: 0104 row: A column: 16
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                  Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Rese
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19 bp DNA linear GSS 02
1M0104A16R Mouse 10kb plasmid UUGCIM library Mus musculus
clone UUGCIM0104A16 R, genomic survey sequence.
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GSS.
                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
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USA
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/note="Vector: pWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
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/lab_host="E. Coli strain XL10-Gold,
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/clone="UUGC1M0016E23"
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                                                                                                                                                                                                                                                                                                                                                                                                                                     19 bp DNA linear GSS 03-OCT-200100157609R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0157G09 R, genomic survey sequence.
                                                                                                                                                                                                    1 (bases 1 to 19)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                             University of Utah
                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                         Mus musculus
                                                                                                                         Contact: Robert B.
                                                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                  AZ394192.1 GI:10509264
GSS.
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Location/Qualifiers
801 585 5606
801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli X110-Gold (Stratagene) cells and selected for ampicillin resistance."
                                         USA
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                                                          Biomedical Polymers Research Bldg.,
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/sex="Male"
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/strain="C57BL/6J"
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                   JOURNAL COMMENT
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US-09-966-880A-8 (1-198) x AZ394192 (1-19)
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                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. (bases 1 to 19)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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High quality sequence stop: 19.
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Insert Length: 10000 Std Erro
Plate: 0157 row: G column: 0
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ418201 19 bp DNA linear GSS 03 190194M12F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0194M12 F, genomic survey sequence.
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                                                                                             plasmid inserts
                                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                     Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                             Eukaryota; Metazoa; C
Mammalia; Eutheria; F
1 (bases 1 to 19)
Dunn,D., Aoyagi,A., F
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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AZ441188.1 GI:10565117
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Insert Length: 10000 Std Error:
Plate: 0194 row: M column: 12
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                Mus musculus
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Class: plasmid ends
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Rm. 308, Biomedical
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/strain="C57BL/6J"
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    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                         Mus musculus
                                                                                                                                                                    AZ447414.1
GSS.
                                                                                                                                                                                                                                                                           AZ447414 19 bp DNA linear GSS 04-OCT-20 1M0244L06R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0244L06 R, genomic survey sequence.
                                                                                                                       Mus musculus (house mouse)
                                                                                                                                                                                                                                              AZ447414
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Insert Length: 10000 Std Erro
Plate: 0232 row: O column: 0
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Fax: 801 585 7177
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gh quality sequence stop: 19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNP42 (gi 4732114 |gb | AF12972.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone="UUGC1M0232004"
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/strain="C57BL/6J"
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AZ478491 19 bp DNA linear GSS 04-OCT-200 IM0298903R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0298P03 R, genomic survey sequence.

AZ478491 AZ478491.1 GI:10637389
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Class: plasmid ends
High quality sequence stop: 19.
Location/Qualifiers
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Unpublished (2000)
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Fax: 801 585 7177
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                                                                                                                                                                                                 GGAAGGAGGGAG 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared NNA was blunt end-repaired with 74 DNA polymerase and 74 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWp42 (g1|4732194|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Length: 10000 Std Error: 0.00
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strain="C57BL/6J"
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AUTHORS
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27 ThrTyrLeuCys 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0298 row: P column: 03
Seq primer: CACACAGGAAACAGCTATGACC
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University of Utah Genome Center
University of Utah
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1 (bases 1 to 19)
Dunn, D., Aoyagi, A., Barber, M.,
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Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 308,
                                                                                                                                                                                                                                                                                                                   (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWH42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:10090"
/clone="UUGC1M0298P03"
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                                                                                                                     100.00%
100.00%
2.02%
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4.00
                                                 (1-19)
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Matches:
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	-880A-8 (1-198) x AZ498063 (1-19)	US-09-966-
) }
	Percent Similarity: 100.00% Conservative: 0 Best Local Similarity: 100.00% Mismatches: 0 Query Match: 2.02% Indels: 0 DB: 28 Gaps: 0	Percent Simi Best Local S Query Match: DB:
	4.48e+05 Length:	Alignment Pred. No.: Score:
	polynucleotide kinase. Adaptor oilgonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi 4732114 gb AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."	ORIGIN
	crain XL10- cb plasmid nv; Purific ale) was oh Resources/c resources/c sheared by constant d with T4	
	/organism="mus musculus" /mol_type="genomic DNA" /strain="C57BL/6J" /db_xref="taxon:10090" /clone="UUGC1M0335J08" /sex="Male"	
	Location/Qualii	FEATURES source
·	Fax: 801 585 7177 Fax: 801 585 7177 Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: 0.00 Plate: 0335 row: J column: 08 Seq primer: CCTTGTAAAACGACGCCAGT Class: plasmid ends High quality sequence stop: 19.	
·····	University of Utah University of Utah Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 84112, USA 701-801 FOR FORC	
	prasmid inserts Unpublished (2000) Contact: Robert B. Weiss	JOURNAL COMMENT
	cre, S., Mahmoud, M., Meenen, E., Pederse, M., Rose, R., Stokes, R., Tingey, A., vo. and Wright, D. Weises, R. ome scaffolding with paired end reads	TITLE
-	.5.	SOURCE ORGANISM REFERENCE AUTHORS
	AZ498063 1M0335J08F MOCLONE UUGC1M0 AZ498063 AZ498063.1 G	LOCUS DEFINITION ACCESSION VERSION KEYWORDS
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4 A S A S A S A S

105 SerLeuArgIle 108

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VERSION
KEYWORDS
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Pred. No.:
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AUTHORS
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AZ582154
  Percent Similarity:
Best Local Similarity:
Query Match:
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Insert Length: 10000 Std Error: 0.00
Plate: 0374 row: C column: 19
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 19.
Location/Qualifiers
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Fax: 801 585 7177
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AZ582154.1 GI:11700755
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                                                                                                                                                                                             /lab host="E. Coli strain X110-Gold, T1-resistant, F-"
//lab host="E. Coli strain X110-Gold, T1-resistant, F-"
//clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42ny, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWP42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db xref="taxon:10090"
/clone="UUGC1M0374C19"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sex="Male"
  4.00
100.00%
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2.02%
                                                                                                     4.48e+05
Conservative: Mismatches: Indels:
  4000
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RESULT 23
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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Insert Length: 10000 Std Error: 0.00
Plate: 0407 row: C column: 19
Seq primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
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Contact: Robert B.
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            10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwd42 (gil4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                               Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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/clone lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'db_xref="taxon:10090"
'clone="UUGC1M0407C19"
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strain="C57BL/6J"
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Location/Qualifiers
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1M0407C15R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC1M0407C15 R, genomic survey sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
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Mammalia; Butheria; Rodentia;
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil 4732114 [gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  clone="UUGC1M0407C15"
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High quality sequence stop: 19.
"https://publiciers
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Plate: 0448 row: M column: 12
Seq primer: CGTTGTAAAACGACGGCCAGT
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1M0448M12F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0448M12 F, genomic survey sequence.
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blint end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative
                                                                                                                                                                                                            /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                 sex="Male"
                                                                                                                                                                                                                                                                                                                                                       clone="UUGC1M0448M12"
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                                                                                                                                                                                                                                                                                                                                                       source
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                                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop: 19.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
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19 bp DNA linear GSS 14
1M0540I06R Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0540I06 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plate: 0540 row: I column: 06
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
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GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                           Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Smail: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plasmid inserts
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
                                                                                           /lab_host="E. Coli strain Xi10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCNM library"
/note="Vector: PWD42nv, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                     clone="UUGC1M0540I06"
                                                                                                                                                                                                                                                           db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                 /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                               organism="Mus musculus"/
                                                                                                                                                                                                                    Bex="Male"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUTHORS
                                                                                                                                                                                                                                          source
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                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0026 row: M column: 20
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
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AZ784061.1 GI:12919427
GSS.
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2M0026M20F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0026M20 F, genomic survey sequence.
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2000)
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
/lab_host="B. Coli strain XL10-Gold, Tl-resistant, /Clone_lib="Mouse 10kb plasmid UUGC1M library" /note="Wector: PWD42nv; Purified genomic DNA from Mouseulus C57BL/6J (male) was obtained from the Jac
                                                                                                                                              /mol_type="genomic DNA"
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                                                                                                                       clone="UUGC2M0026M20"
                                                                                                                                                                                                                   organism="Mus musculus"
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                           from M.
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SOURCE

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AUTHORS

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/db_xref="taxon:10090" /clone="UUGC2M0056J18"

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2M0056J18R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0056J1B R, genomic survey sequence.
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                                                                                                                          row: J column: 18
Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality -
                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 19)
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GSS.
Mus musculus (house mouse)
                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0056 row: J column: 1
                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
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/organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                           Location/Qualifiers
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2M0117N04F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0117N04 F, genomic survey secmence
                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0117 row: N column: 04
                                                                                                                                                                                                                                                                                                                                        Contact: Robert B. Weiss University of Utah Genome University of Utah
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                                                                               Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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Mus musculus
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801 585 7177
quality sequence stop: 19
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(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pND42 (gi |4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                         USA
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RESULT 30
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                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0174 row: M column: 11
                                                                                                                                                         Contact: Robert B. Weiss University of Utah Genome University of Utah
                                                                                                                                                                                                                                 Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ864551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2M0174M11F Mouse 10kb plasmid \overline{U}UGC1M library Mus musculus genomic clone \overline{U}UGC2M0174M11 F, genomic survey sequence.
                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                                                                          Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                               Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                   USA
                                                                                                                                      Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Laboratory Mouse DNA Resource
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/mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                             Genome Research Group
National Institute of Radiological Sciences
9-1, Anagawa-4-chome. Inage-ku, Chiba, Chiba 263-8555, Japan
                                                                                                                                                                          1 (bases 1 to 20)
Morimyo, M. and Mita, K.
Identification of expressed
                                                                                                                                                                                                                                                                                    Schizosaccharomyces pombe (fission yeast)
Schizosaccharomyces pombe
Eukaryota; Pungi; Ascomycota; Schizosaccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends High quality sequence stop: 19.
               Email: morimyo@nirs.go.jp.
Location/Qualifiers
                                                                                                                    Contact: Mitsuoki Morimyo
                                                                                                                                                                                                                                                                                                                                                                          AU013258
AU013258.1 GI:3368049
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quality sequence stop: 19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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strain="C57BL/6J"
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Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata;
Mammalla; Eutheria; Rodentia;
1 (bases 1 to 20)
Kato,K. and Matoba,R.
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                                                                                                                                                                                                                                                                                                            URL:http://love2.aist-nara.ac.jp/BED/index.html.
                                                                                                                                                                                                                                                                                                                                                  Tel: 81-743-72-5581
Fax: 81-743-72-5589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                 Email: kkato@bs.aist-nara.ac.jp
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                                                                                                                                                           /tissue_type="brain"
/clone_lib="3'-directed mouse cDNA library"
                                                                                                                                                                                                                    organism="Mus musculus"
/mol_type="mRNA"
/db_xref="taxon:10090"
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/sex="h minus"
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/mol_type="mdxA"
/strain="972"
/db_xref="taxon:4896"
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                                  Mismatches:
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и 40000
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Percent Similarity:
Best Local Similarity:
Query Match:
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BX558127
                                                                                                             RESULT 34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
All clones with suffix qlc are i
end of the cDNA all plc reads as
20 bp mRNA linear EST 01-AUG-2003 D09C05 E05 05.abl cDNA Peking library 2, 4 day SCN3 Glycine max cDNA clone D09C05 5', mRNA sequence.
CA851019
CA851019.1 GI:33387612
                                                                                                                                                                    73 GlyArgCysTyr 76
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The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neopteara; Endopterygota; Diptera; Brachycera; Muscomorpha; Hippoboscoidea; Glossinidae; Glossina.

1 (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BX558127 Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans cDNA clone Tse36g09_glc, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Berriman,M., Hamilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S. and Hall,N. Adult midgut expressed sequence tags from the tsetse fly Glossina morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Glossina morsitans
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Genome Biol. 4 (10), R63 (2003)
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                                                                                                                                           GGGAGGTGTTAT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Glossina morsitans
/mol_type="mRNA"
/sub_species="morsitans"
                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="Tse36g09_q1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
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                                                                                                                                                                                                                                                                                                                                                                                         note="country:
.brucei"
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CF322764/c
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AUTHORS
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Best Local Similarity:
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        Yongin, Kyeonggi, K
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                       Oryza sativa
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US-09-966-880A-8 (1-198) x CA851019
                                                                                     Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzae; Oryza.

1 (bases 1 to 20)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               HDN--02-A02.g1 OBHDAC1-overexpressing
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Alkharouf, N. W., Khan, R. and Matthews, B.F.
Analysis of expressed sequence tags from
infected by the soybean cyst nematode
Unpublished (2002)
                    Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                        Contact: Nahm B.H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Soybean Genomics and Improvement Laboratory (SGIL) US Department of Agriculture (USDA), ARS, PSI Bldg.006, Rm 118, 10300 Baltimore Ave., Beltsville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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Glycine max (soybean)
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alkharon@ba.ars.usda.gov.
Location/Qualifiers
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/dev.stage="Seedlings"
/dev.stage="Seedlings"
/clome_lib="cDNA_peking_library_2, 4 day_SCN3"
/note="Vector: pBluescript_SK-; cDNA_clones from mRNA_extracted_from_Peking_roots_2 and 4 days_past_invasion
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|mol_type="mRNA"
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'cultivar="Peking"
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gl OsHDAC1-overexpressing transgenic rice lambda ph
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice BSTs

Unpublished (2003)
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CF327699.1 GI:33803647
                                                                                                                                                                                                                                                                                                                                      of Bioscience and Bioinformatics,
                                                                                                                                                                                                                                                                                                                                                                            Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oryza sativa
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82 31 321 6355
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Location/Qualifiers
                                                                                                                                                                                                                                                          bhnahm@ggbio.com,
                                                                                                                                                                                                                                                                                                                 Kyeonggi,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: XhoI; cDNA was inserted into lambda UnT-ZAP XR vector at 5' end with EcoRI and 3' end with KhoI site. mRNA was derived from rice Histone Deacetylase overexpression
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/dev_stage="proliferated callus on 2N6 media for 30 days"
/lab_host="E.coli DH108"
/lab_host="E.coli DH108"
/clone lib="Rice callus plasmid cDNA library (NACL)"
/note="Vector: pCR4-TOPO; Site 1: EcoRI; mRNA was capped
with oligoribonucleotides and then used as templates for
RT-PCR."
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/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                        /cultivar="Nackdong"
/db_xref="taxon:4530"
/clone="NACL--02-E17"
                                                                                                                                                                                   /organism="Oryza sativa"
/mol_type="mRNA"
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dev_stage="proliferated callus on 2N6 media for 2 weeks"
                                                                                                        tissue_type="callus"
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AZ303578/c
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US-09-966-880A-8 (1-198) x CF327699 (1-20)
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1M0003H07F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0003H07 F, genomic survey sequence.
AZ303578
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Location/Qualifiers
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Insert Length: 10000 Std Error:
Plate: 0003 row: H column: 07
Seq primer: CGTTGTAAAACGACGGCCAGT
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi14732114|gb|AF119072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                                                                                                                                            /lab host="E. Coli strain XL10-Gold, T1-resistant, F-" /clome_1b="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWP42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                sex="Male"
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'clone="UUGC1M0003H07"
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AZ307763/c
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb
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1M0010F06F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGCIM0010F06 F, genomic survey sequence.
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Insert Length: 10000 Std Error:
Plate: 0010 row: F column: 06
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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                                                                                                                                                                                                                                                                                      /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"/note="Vector: PWD42nv; Purified_genomic_DNA_from_M.musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtai
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Mus musculus"
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RESULT 39
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                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606

Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0059 row: D column: 2:
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1M0059D21R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0059D21 R, genomic survey sequence
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Seq primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B.
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/lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: PWD42rv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg.,
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                                                                                                                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                        location/Qualifiers
                                                                                             sex="Male"
                                                                                                                    clone="UUGC1M0059D21"
                                                                                                                                          db_xref="taxon:10090"
                                                                                                                                                                                                           organism="Mus musculus"
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Laboratory Mouse DNA Resource

http://www.jax.org/resources/documents/dnares/). The DNA

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REFERENCE
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Best Local Similarity:
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1M0084F13R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0084F13 R, genomic survey sequence
AZ348201
                                                                                                                                                                                                     Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0084 row: F column: 13
                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B. Weiss
University of Utah Genome
University of Utah
Rm. 308, Biomedical Polyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,B., Pedersen,T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus (bases 1 to 20)
                                                                                                                                                           High quality sequence stop: 20.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                           USA
                                         /organism="Mus musculus"
/mol type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                 clone="UUGC1M0084F13"
sex="Male"
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_host="E. Coli strain XL10-Gold, T1-resistant,

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RESULT 41 AZ387854/c SOURCE ORGANISM Percent Similarity: Best Local Similarity: Alignment Scores: Pred. No.: REFERENCE US-09-966-880A-8 (1-198) x AZ348201 (1-20) Query Match: DEFINITION ACCESSION AUTHORS 125 GlyLeuArgArg 128 Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. 20 bp DNA linear GSS 02-OCT-200 1M0147K24R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0147K24 R, genomic survey sequence տ Tel: 801 585 5606 Fax: 801 585 7177 Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: 0.00 Plate: 0147 row: K column: 24 University of Utah (University of Utah Rm. 308, Biomedical Unpublished (2000) Contact: Robert B. Mus musculus Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends AZ387854 Mus musculus (house mouse) AZ387854.1 GI:10501562 GGACTGAGGAGA 16 quality sequence stop: 20.
Location/Qualifiers was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with TA DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel elettrophoresis. Vector DNA was prepared from a derivative of putative agarose with adaptore complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent 8. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance." USA /clone_lib="Mouse 10kb plasmid UUGC1M library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource /organism="Mus musculus" /mol_type="genomic DNA" (http://www.jax.org/resources/documents/dnares/). 100.00% 100.00% 2.02% 4.00 4.73e+05 Genome Center Weiss Polymers Research Bldg., Length: Matches: Indels: Mismatches: Conservative: ×40000 20 2030 E., GSS 02-OCT-2000 Ş

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KEYWORDS

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US-09-966-880A-8 (1-198) x AZ387854 (1-20)
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                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 20)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0.66 row: C column: 11
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 bp DNA linear GSS 03-OCT-20.
1M0166C11R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0166C11 R, genomic survey secuence
AZ400367
                                                                                                                                                                                                                                                                                                   Unpublished (2000)
                                                                                                                                                                                                                                                                                                                         plasmid inserts
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus
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                                                                                                                                                                                          308,
                                                                                                                                                                   USA
                                                                                                                                                                                    Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Laboratory Mouse DNA Resource
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/db_xref="taxon:10090"
/clone="UUGC1M0147K24"
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Pred. No.:

RESULT 42

rocus

COMMENT

plasmid ends

JOURNAL TITLE REFERENCE

AUTHORS

SOURCE

ORGANISM

KEYWORDS VERSION ACCESSION DEFINITION AZ400362

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RESULT 43
AZ408559
LOCUS
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AUTHORS
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.

Nouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ408559 20 bp DNA linear GSS 03 1M0179K14R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0179K14 R, genomic survey sequence.
                                                                                                                               University of Utah University of Utah Rm. 308, Biomedical
                                                                                                                                                                                                                                                                                            Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                         plasmid inserts
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#112, USA
#1: 801 585 5606
#X: 801 585 7177
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                               Polymers Research Bldg.,
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                            COMMENT
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                                                      JOURNAL
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                                                                                                                                                                                                      AUTHORS
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                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse,Whole genome scaffolding with paired end reads from 10kb Nouse, whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                       AZ486007 20 bp DNA linear GSS 05-OCT-200 1M0313E17R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0313E17 R, genomic survey sequence.
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                    AZ486007.1 GI:10652355
GSS.
                                                                          plasmid inserts
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Location/Qualifiers
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C578L/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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/clone_lib="Mouse 10kb plasmid_UUGCIM library"
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; 1 (Dases 1 to 20)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                           20 bp DNA linear GSS 05-OCT-200 1M0319H15R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0319H15 R, genomic survey secuence
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Insert Length: 10000 Std Error: (
Plate: 0313 row: E column: 17
Seq primer: CACACAGGAACAGCTATGACC
Class: plasmid ends
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Fax: 801 585 7177
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Mus musculus
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AZ489135.1 GI:10658589
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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/strain="C57BL/6J"
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Insert Length: 10000 Std Error: 0.00
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  Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Fax: 801 585 7177
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/strain="C57BL/6J"
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Sciurognathi; Muridae; Murinae; Mus.
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20 bp DNA linear GSS 14-DEC-20 1M0520A16R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0520A16 R, genomic survey sequence.
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Insert Length: 10000 Extor: (
Plate: 0484 row: J column: 01
Seq primer: CACACAGGAAACAGCTATGACC
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Fax: 801 585 7177
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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Mouse whole genome scaffolding with paired end reads from 10kb
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Mus musculus (house mouse)

Mus musculus
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Fax: 801 585 7177
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Insert Length: 10000 Std Error:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/note="Vector: PWP02nv, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
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/mol_type="genomic DNA"
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Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads
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University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20
84112, USA
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Insert Length: 10000 Std Error: 0.00
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                 http://www.jax.org/resources/documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.05 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of phasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored west and and transformed into
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                                                                                                                                                                                                                           adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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University of Utah Ge
University of Utah
Rm. 308, Biomedical F
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R. Mouse, whole genome scaffolding with paired end reads from 10kb Nouse, whole genome scaffolding with paired end reads from 10kb
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1M0583M24R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0583M24 R, genomic survey sequence
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Insert Length: 10000 Std Error: 0.00
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Fax: 801 585 7177
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/mol_type="genomic DNA"
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
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Insert Length: 10000 Std Erro
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Fax: 801 585 7177
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2M0008K09R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
                             was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with TA DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of phasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
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Alignment Scores:

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RESULT 51
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AZ776071.1 GI:12903267
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20 bp DNA linear GSS 16-FEB-20 200009L24F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0009L24 F, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20)
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Fax: 801 585 7177
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University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (house mouse)
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                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.05 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDM2 (gil4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: FWD42nv; Purified genomic DNA from M. musculus C578L/6U (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from AZ810573 20 bp DNA linear GSS 20-FEB-20 2M0076K11F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0076K11 F, genomic survey sequence. High quality sequence stop: 20. Location/Qualifiers Plate: 0076 row: K column: 11 Seq primer: CGTTGTAAAACGACGGCCAGT Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: Plate: 0076 row: K column: 11 Contact: Robert B. Weiss University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Research Bldg., Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (basee 1 to 20) Class: plasmid ends Tel: 801 585 5606 Fax: 801 585 7177 Unpublished (2000) plasmid inserts Mus musculus Mus musculus (house mouse) AZ810573.1 GI:12977957 USA /organism="Mus musculus"
/mol_type="genomic DNA" 0.00 Duval, B., Hamil, C., 20 S. 2030 E., GSS 20-FEB-2001 Murinae; Mus. SLC,

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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative

/lab_host="E. Coli strain XI10-Gold, T1-resistant, F-"
/clome lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource

'db_xref="taxon:10090" 'clone="UUGC2M0076K11" strain="C57BL/6J"

Bex="Male"

of pWD42 (gi |4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells

selected for ampicillin resistance."

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RESULT 53
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                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0113 row: M column: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1. (bases 1 to 20)
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Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pW042nv, Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                 clone="UUGC2M0113M11"
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                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="C57BL/6J"
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polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XX10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Unpublished (2000)
Contact: Robert B.
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                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0116 row: A column: 09
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                     University of Utah Genome Center University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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200116A09R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
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igh quality sequence stop: 20.
Location/Qualifiers
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/lab_host="B. Coli strain XL10-Gold, T1-resistant, F. /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PW042nv; Purified genomic DNA from M. /note="Vector: PW042nv; Purified genomic DNA from M. /note="Vector: PW042nv; Purified genomic DNA from the Jack."
                                                                                                                                /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                      db xref="taxon:10090"
                                                                                                  clone="UUGC2M0116A09"
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The

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JOURNAL COMMENT
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AZ949545
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                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0213 row: E column: 19
Seq primer: GGTGTAAAACGACGGCCAGT
Class: plasmid ende
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Rese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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Fax: 801 585 7177
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20 bp DNA linear GSS 27-APR-20
2M0213E19F Mouse 10kb plasmid UUGC2M library Mus musculus genomic
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quality sequence stop: 20.
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/mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                 organism="Mus musculus"
                                                                                                           ocation/Qualifiers
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KEYWORDS VERSION

FEATURES

/db_xref="taxon:3847" /clone="807D03"

tissue_type="Roots"

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REFERENCE
AUTHORS
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Best Local Similarity:
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Alkharouf,N.W., Khan,R. and Matthews,B.F.
Analysis of expressed sequence tags from roots of resistant soybean infected by the soybean cyst nematode
Unpublished (2002)
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Glycine max (soybean)

Glycine max

Glycine max

Glycine max

Streptophyta; Embryophyta; Tracheophyta;

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;

rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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B07D03 seq cDNA Peking library 12hr SCN3 Glycine max cDNA clone
B07D03 5', mRNA sequence.
CA853358
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                                                                                                                                                      Tel: 301 504 5750 Fax: 301 504 5728
                                                                                                                                                                                                                                  Bldg.006, Rm
                                                                                                                                                                                                                                                           Soybean Genomics and Improvement Laboratory (SGIL)
US Department of Agriculture (USDA), ARS, PSI
                                                                                                                                                                                                                                                                                                                    Contact: Alkharouf, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CA853358.1 GI:33390151
                                                                                                  alkharon@ba.ars.usda.gov.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gi|473214|gb|AFT29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
/organism="Glycine max"
/mol_type="mRNA"
/cultivar="Peking"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  musculus C57BL/6J (female) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="B. coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'sex="Female"
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4.00
                                                                                                                                                                                                                                     118,
                                                                                                                                                                                                                                  10300 Baltimore Ave., Beltsville, MD 20705-2350,
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                                Alignment Scores: Pred. No.:
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HSM004456/c
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                                                                              Sequence 10
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                                                                                                                                                                                                                                                                                                   This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, He Berlin-Charlottenburg, GERMANY; Email: clone@
                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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12-MAR-1999
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                               Clone from S. Wiemann, sequenced by Qiagen within the cDNA sequencing consortium of the German Genome Project
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/clone_lib="cDNA Peking library 12hr SCN3"
/note="Vector: pBluescript SK-; cDNA clones from mRNA
extracted from roots of soybean cv. Peking 12 hrs after
infection by SCN race 3. These are cloned in pBluescript
SK- phagemid. "
                                                                                                         /clone_lib="434 (synonym: htes3). Vector
DH10B; sites Not! + Sal!"
/dev_stage="adul!"
/tissue_type="testis"
                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                         organism="Homo sapiens"
|clone="DKFZp434L1012"
                                                                                                                                                                                                      /db_xref="taxon:9606"
/mol_type="mRNA"
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Last updated, Version
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                                                                            2 T; 0 other;
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Matches:
Conservative:
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RESULT 59
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RESULT 58
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Query Match:
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                                                                                              US-09-966-880A-8 (1-198)
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Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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12-MAR-1999
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Please contact the RZPD: Ressourcenzentrum, He
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                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:9606"
/mol_rype="mRNA"
/organism="Homo sapiens"
/clone="DKFZp434J1415"
/clone_lib="434 (synonym: hi
DH10B; sites Noti + Sall"
/dev_stage="adult"
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                                                                                                                                                                                                                                                                                                                                                                                           tissue_type="testis"
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Last updated, Version
                                                                                              HSM005384
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best Local Similarity:
Query Match:
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CF323895
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                                                                REFERENCE
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                                                AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                            10 GGCCGACGG
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1 (bases 1 to 10)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
                                                                                                                                                                                                                                                                   CF323895 10 bp mRNA linear EST 18-AU HDN--05-A22.gl OsHDAC1-overexpressing transgenic rice lambda cDNA library II (HDN) Oryza sativa cDNA clone HDN--05-A22, mJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Nahm B.H.
Genomics and Genetics Institute, Greendene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Lazge-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
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                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                Oryza sativa
                                                                                                                                                                                                        CF323895.1 GI:33796055
                                                                                                                                                                                                                                              sequence.
                                                                                                                                                              Oryza sativa
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: pCR4-TOPO; Site_1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cDNA library (HD)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="B.coli DH10B"
/clone_lib="0sHDAC1-overexpressing transgenic rice plasmid
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'cultivar="Nackdong"
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Matches:
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                                           Shin, Y.C.,
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CF339022/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOURCE
ORGANISM
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AUTHORS
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JOURNAL
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)

Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Oryza sativa
Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 bp mRNA linear ERCL1--03-I17.gl Regenerated callus lambda phage cDNA Oryza sativa cDNA clone RCL1--03-I17, mRNA sequence.
                                                                                                                                                            Genòmics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193
Fax: 82 31 321 6355
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CF339022.1 GI:33826427
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                                                                                                                       bhnahm@ggbio.com, bhna
Location/Qualifiers
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/mol type="mRNA"
/cultivar="Nackdong"
/db xref="taxon:4530"
/clone="HDN--05-A22"
                       /organism="Oryza sativa"
/mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:4530"
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                                                                                                                                              bhnahm@bio.myongji.ac.kr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length:
Matches:
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Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: XhoI; cDNA was inserted into lambda Uni-ZAP XR vector at 5' end with EcoRI and 3' end with XhoI site. mRNA was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="OsHDAC1-overexpressing transgenic rice lambda
phage cDNA library II (HDN)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  derived from rice Histone Deacetylase overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /dev stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="B.coli SOLR"
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                                                                                                                                                                                                                                                                                     0000#6
                          EST 18-AUG-2003
library (RCL1)
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/clone="RCL1--03-I17" /tlssue_type="callus" /dev_stage="proliferated callus on 2N6 media for 30 days" /lab_host="B.coli SOLR"

/clone_lib="Regenerated callus lambda phage cDNA library (RCL1) "

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RESULT 62
HSM008167/c
ID HSM008167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; ver
Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo
                                                                                                                                                                                                                                                                                           Clone from S. Wiemann, sequenced by LMU within the cDNA sequencing consortium of the German Genome Project
                                                                                                                                                                                                                                                                                                                                                                                              Blum
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12-MAR-1999
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                                                                                                                                                                                                                             Berlin-Charlottenburg,
                                                                                                                                                                                                                                            No s1 sequence available
This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Ho
 Sequence 11
                                                                                                                                                                                                                                                                                                                                                               Submitted
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   BP;
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/clone="DKFZp434N0723"
/clone="DKFZp434N0723"
/clone="DKFZp434N0723"
/sib="A34" (synonym: h
DH10B; sites Not! + Sal!"
/dev_stage="adult"
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 2 A;
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                                                                                                                                                                                               Location/Qualifiers
                                                                                                                             /db_xref="taxon:9606"
/mol_type="mRNA"
                               'tissue_type="testis"
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 2 C;
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                                                                                                                                                                                                                             D: Ressourcenzentrum, Heubnerweg GERMANY; Email: clone@rzpd.de
                                                                                                                                                                                                                                                                                                                                                                                         Mewes W., Gassenhuber J.,
4 G;
 3 T; 0 other;
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REFERENCE
AUTHORS
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BG896271
                                                                                        Best Local Similarity:
Query Match:
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Best Local Similarity:
                                                        US-09-966-880A-8 (1-198) x BG896271
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HOA28-1-G6 HOA (Human Osteoarthritic Cartilage) Homo
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                                                                                                                                                                                                                                                                                                                                                                                      Seq primer:
                                                                                                                                                                                                                                                                                                                                                                                                                                     709 Swedeland Road, Tel: 610-270-7245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and osteoarthric cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 11)
                                                                                                                                                                                                                                                                                                                                                                                                      Email: sanjay_kumar-1@gsk.com
                                                                                                                                                                                                                                                                                                                                                                                                                        Fax: 610-270-5598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GlaxoSmithKline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Sanjay Kumar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kumar, S., Connor, J.R., Dodds, R.A., Halsey, W., Van Horn, M., Mao, J.,
Sathe, G., Mui, P., Agarwal, P., Badger, A.M., Lee, J.C., Gowen, M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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GCCAGACTC 11
                                                                                                                                                                                                                                /lab hoste"E.coli DH10 B"
/clone lib="HOA (Human Osteoarthritic Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
Directional"
                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                            tissue_type="cartilage"
                                                                                                                                                                                                                                                                                                                                                                           location/Qualifiers
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                                                          (1-11)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Box 1539, King of Prussia,
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Matches:
Conservative:
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Matches:
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                                                           1 (bases 1 to 11)
Turkewitz,A.P., Karrer,K.M.,
Frankel,J. and Klobutcher,L.
                                                                                                              Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota, Alveolata, Ciliophora, Oligohymenophorea,
Hymenostomatida, Tetrahymenina, Tetrahymena.
                                                                                                                                                                                                                                           BM395226 11 bp mRNA linear EST 17 50072-2-8-B04.f.2 Chilcoat/Turkewitz cDNA (large fraction)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and osteoarthritic cartilage cDNA libraries
Osteoarthr. Cartil. 9 (7), 641-653 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 11)

Kumar,S., Connor,J.R., Dodds,R.A., Halsey,W., Van Horn,M., Mao,J.
Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen,M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BG927412
BG927412.1 GI:14321935
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HNC1-1-G11.R HNC
Unpublished (2002)
Contact: Turkewitz AP
Molecular Genetics and
                                                                                                                                                                                                                Tetrahymena thermophila
BM395226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                 from Tetrahymena thermophila, strain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"

/mol type="mRNA"

/db Xref="reaxon:9606"

/tissue type="cartilage"

/lab host="B.coli DH10 B"

/clone_lib="HNC (Human Normal Cartilage)"

/note="Vector: psport I; Site_1: SalI; Site_2: NotI;

Directional"
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                                                                             Karrer, K.M., Jahn, C.,
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and Cell Biology
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                                                                                 Kirk, K.E.,
                                               growing cells
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920 E. 58th Street, Chi
Tel: 773 702 4374
Fax: 773 702 3172
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                                                                                                                                                                      Seq
                                                                                                                                                                                                      University of Chicago
920 E. 58th Street, Chicago,
Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11 bp mRNA linear EST 17 5009-0-15-C03.t.1 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                Unpublished (2002)
Contact: Turkewitz AP
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Turkewitz,A.P., Karrer,K.M.,
Frankel,J. and Klobutcher,L.
                                                                                                                                                                                                                                                                                                                                                                                        Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
                                                                                                                                                                                                                                                                Molecular Genetics and Cell Biology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       il: apturkew@midway.uchicago.
primer: T3.
                                                                                                                                                                        primer: T3
                                                                                                                                                                                      apturkew@midway.uchicago.edu
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)" /note="Vector: BlueScript2 SK+; Details on library preparation can be found in Chilcoat and Turkewitz (2001) Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                         /organism="Tetrahymena
/mol_type="mRNA"
/strain="CU428.1"
                                                                                                                                                        Location/Qualifiers
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/strain="CU428.1"
/db_xref="taxon:5911"
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Length: Matches: Conservative:

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eudicots;

Tracheophyta;

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RESULT 67
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Line Meg 10, 50829 Koeln, Germany
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H. and Radelof,U.
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BQ587100.1 GI:26116682
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1 (bases 1 to 11)
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                                                                                                                                                                                          Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database:http://gabi.rzpd.de"
                                                                                                                                                                                                                                                                                                                                        /clone lib="MPIZ-ADIS-024-leaf"
/note="Vector: pCMVSPORT6; Site 1: SalI; Site 2: NotI;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
                                                                                                                                                                                                                                                                               SP6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7;
                                                                                                                                                                                                                                                                                                                      b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="leaf"
/lab_host="EMDH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Socation/Qualifiers
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|mol_type="mRNA"
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Mismatches:
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Pred. No.:
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BQ595495
LOCUS
                                                                                                         Query Match:
                                                                                                                          Percent Similarity:
Best Local Similarity:
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ORGANISM
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                                       US-09-966-880A-8 (1-198) x BQ595495 (1-11)
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MEDLINE
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  181 LeuProLeu 183
                                                                                                                                                                                                               Scores:
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eu
Caryophyllales; Amaranthaceae; Beta.
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Plate: 22 row: O column: 14
Seg primer: SP6; CATACGATTTAGGTGACACTATAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11 bp mRNA linear EST 06-DEC-2002
E012591-024-022-014-SP6 MPIZ-ADIS-024-developing root Beta vulgaris
ED02525 Clone 024-022-014 5-PRIME, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Weisshaar B
ADIS DNA core facility at MPIZ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: weisshaa@mpiz-koeln.mpg.de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Max-Planck-Institute for Plant Breeding Research
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                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="MPIZ-ADIS-024-developing root"
/note="Vector: pCMVSPORT6; Site 1: Sal1; Site 2: Not1;
cDNA library from sugar beet, library provided by KMS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                          Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinate Prof. Christian Jung; Sequence submission managed by
                                                                                                                                                                                                                                                                                                                                                         SP6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7;
                                                                                                                                                                                                                                                                        RZPD/GABI-Primary database: http://gabi.rzpd.de'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="developing root"
/lab_host="EMDH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="GABI:191359"
/db_xref="taxon:161934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="mxxx"
/cultivar="KWS2320 (double haploid, monogerm breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Beta vulgaris"
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AUTHORS
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CF543159
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Best Local Similarity:
Query Match:
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                                                                                                                                                                      CF543159 11 bp mRNA linear EST 22-SEP-2003 S014678-024-030-006-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Amaranthaceae; Beta.
                                                        Beta vulgaris
Beta vulgaris
                                                                                                                                      024-030-006 5-PRIME,
CF543159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.,
of Bioscience and Bioinformatics, MyongJi University
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Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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                                                                                                                  CF543159.1 GI:34891599
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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                                                                                                                                                                                                                                                                            AGTGCCACC
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/db_xref="taxon:4530"
/clone="RCL1--03-K22"
/tissue_type="callus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="Vector: pBluescript SK(+); Site 1: SatI; Site 2: XhOI; CDNA was inserted into lamda Uni-ZAP XR vector at 5' end with SatI and 3' end with XhOI site. Callus was induced on 2N6 media for 30 days and cultured for 36hrs on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   regenerated
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/lab_host="E.coli SOLR"
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|mol_type="mRNA"
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BH129987
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                                                                                                                                                                                                                                                     BH129987 11 bp DNA linear G-6e3.f Maize Random Small-insert Genomic Library clone G-6e3 both, genomic survey sequence.
  Abundance,
elements in
               1 (bases 1 to 11)
Meyers,B.C., Tingey,S.V. and Morgante,M.
Abundance, distribution and transcriptional
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Max-Planck-Institute for Plant Breeding Research
Max-Planck-Institute for Plant Breeding Research
Cort-Con-Linne Weg 10, 50829 Koeln, Germany
                                                                               Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoldeae; Andropogoneae; Zea.
                                                                                                                                                Zea mays
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                                                                                                                                                                                                                                    BH129987
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Insert Length: 11 Std Error:
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in the maize genome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database:http://gabi.rzpd.de"
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/clone="024-030-006"
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REFERENCE
AUTHORS
TITLE
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BH129987/c
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                Dupont Genomics
PO Box 6104, Newark, 1
PO1: 302 631 2638
Fax: 302 631 2607
                                                                                                                                    clade; rannown.

1 (bases 1 to 11)

1 (bases 1 to 11)

Meyers,B.C., Tingey,S.V. and Morgante,M.

Abundance, distribution and transcriptional

elements in the maize genome

elements in the maize Genome

11 (10), 1660-1676 (2001)
                                                                                                                                                                                                                                                                                                                                                                               BH129987 11 bp DNA linear GSS 23-JUL-200: G-6e3.f Maize Random Small-insert Genomic Library Zea mays genomic clone G-6e3 both, genomic survey sequence.
                                                                                       Suite 200
                                                                                                        Contact: Morgante
                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                      SSD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: Michele.morgante@usa.dupont.com
Sequences were trimmed to include only high quality bases; forward
and reverse reads were assembled when significant overlaps were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PO Box 6104, Newark,
Tel: 302 631 2638
Fax: 302 631 2607
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PO Box 6104, Ne
                                                                                                                                                                                                                                                                                                  Zea mays
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Michele.morgante@usa.dupont.com
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/cell_type="young leaf"
/dev_stage="young leaf"
/dev_stage="seedling"
/clone_lb="Maize Random Small-insert Genomic Library"
/note="Vector: pCR-Script; Total genomic DNA was
nebulized; ends were polished with Pfu polymerase and the
fragments cloned into pCR-Script."
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strain="B73"
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XX AL04308
XX I2-MAR-
DT 12-MAR-
DT 12-MAR-
CX Homo sa
XX EST; ex
XX Homo sa
CC Eukaryo
CC Euka
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                                                                                                                                               No s1 sequence available

No s1 sequence available

This clone is available at the RZPD in Berlin

The clone is available at the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059

Please contact the RZPD: Ressourcenzentrum, Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                                                                                                                                                                                                                                     Clone from S. Wiemann, sequenced by sequencing consortium of the German
                                                                                                                                                                                                                                                                                                                                                                                              Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Blum H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Class: shotgun.
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/db_xref="taxon:9606"
                                                                                                                     Location/Qualifiers
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/note="Vector: pCR-Script; Total genomic DNA was
nebulized; ends were polished with Pfu polymerase and the
fragments cloned into pCR-Script."
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/tissue_type="leaf"
/cell_type="Young leaf"
/dev_stage="seedling"
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59, Last updated, Version 1)
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Genome Project
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 Best Local Similarity:
Query Match:
DB:
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BG925521
LOCUS
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Best Local Similarity:
Query Match:
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BG925521
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HNC5-1-D3.R HNC (Human Normal Cartilage) Homo sapiens cDNA, mRNA
                                                                                                                                                                                                                                                                                                Tel: 610-270-7245
Fax: 610-270-5598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kumar,S., Connor,J.R., Dodds,R.A., Halsey,W., Van Horn,M.,
Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen.
                                                                                                                                                                                                                                                                                                                         709 Swedeland Road, P.O. Box 1539,
                                                                                                                                                                                                                                                                                                                                            GlaxoSmithKline
                                                                                                                                                                                                                                                                                                                                                                                                            osteoarthritic cartilage cDNA libraries
Osteoarthr. Cartil. 9 (7), 641-653 (2001)
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                                                                                                                                                                                                                                                                                                                                                                     Contact: Sanjay Kumar
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                                                                                                                                                                                                                                                                       sanjay_kumar-l@gsk.com
imer: T7.
                                                                                                                         /tissue_type="cartilage"
/lab_host="R.coli DH10 B"
/clone_lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
Directional"
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DH10B; sites NotI + SalI"
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                                                                                                                                                                                                'db_xref="taxon:9606"
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Lee, J.C., Gowen, M. and
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Eukaryota; Viridiplantae; Streptophyta; Embryo
Spermatophyta; Magnoliophyta; eudicotyledons;
Caryophyllales; Amaranthaceae; Beta.
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E012340-024-010-M01-SP6 MPIZ-ADIS-024-leaf Beta
024-010-M01 5-PRIME, mRNA sequence.
BQ587766
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Herwig,R., Schulz,B., Weisshaar,B., Drungowski,M., Stahl,D., Wruck,W., and Radelof,U.
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                                                                                                                                                                                                                                                                                                                                          /clone_lib="MPIZ-ADIS-024-leaf"
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: NotI;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: NotI;
cDNA library from sugar beet, library provided by KWS
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-NotI, primer sites and
                                                                                                                                                                                                                      Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schmeider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database:http://gabi.rzpd.de"
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Sequencing granted in the context of the GABI-Beet
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/mol_type="mRNA"
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/lab_host="EMDH10B"
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Menze,A., O'Brien,J., Lehrach,H.
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicoryledons; core eudicots;

Caryophyllales; Amaranthaceae; Beta.

1 (bases 1 to 12)

1 (bases 1 to 12)

Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,

Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
                                                                                        24 ArgArgGlu 26
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Contact: Weisshaar B
ADIS DNA core facility at MPIZ
ADIS DNA --- Tratitute for Plant Breeding Research
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E012340-024-010-M01-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone
024-010-M01 5-PRIME, mRNA sequence.
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Seq primer: SP6; CATACGATTTAGGTGACACTATAG
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/note="Vector: pcMVSPORT6; Site 1: Sall; Site 2: Notl;
/note="Vector: pcMVSPORT6; Site 1: Sall; Site 2: Notl;
cDNA library from sugar beet, llbrary provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-Notl, primer sites a
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/lab_host="EMDH10B"
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/clone="024-010-M01"
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|mol_type="mRNA"
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'db_xref="taxon:161934
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EST631493 DSCT Colletotrichum
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Department of Plant Pathology
University of Minnesota
495 Borlaug Hall, 1991 Upper Buford Circle, St. Paul,
Tel: 612 625 1243
Fax: 651 649 5058
Email: debbys@puccini.crl.umn.edu
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EST631493 DSCT Colletotrichum trifolii cDNA clone pDSCT1-51, mRNA
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ESTs from mycelia of Colletotrichum trifolii race 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone lib="NSCT"
/clone lib="NSCT"
/clone lib="NSCT"
/note="Vector: pBluescript SK+; Site_1: EcoRI; Site_2:
/note="Vector: pBluescript SK+; Site_1: EcoRI; Site_2:
/note="Vector: pBluescript SK+; Site_1: EcoRI; Site_2:
/note="Vector: pBluescript SK+; Site_1: EcoRI from polyA+
enriched RNA The cDNA was pigated into Lambda gtll from
Stratagene and packaged using Gigapack packaging extracts.
An aliquot of the amplified library was used to transduce
E. coll Y1090 and phage DNA was purified from a liquid
lysate. The cDNA inserts were gel purified after EcoRI
digestion and ligated into pBluescript SK+. Aliquots of
the ligation were used to transform E. coll DH5alpha which
were plated onto medium with X-gal for selection of
recombinants."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="pDSCT1-51"
/tissue_type="mycelia"
/tissue_type="mycelia"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Colletotrichum trifolii"
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                     12 bp
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Indels:
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    trifolii
                     mRINA
A linear EST 18-JUL-2002 CDNA clone pDSCT1-51, mRNA
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C51419/c
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Query Match:
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BQ750930.1 GI:21906335
EST.
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C51419 Yuji Kohara unpublished
clone yk195ell 3', mRNA sequenc
C51419
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Tel: 612 625 1243
Fax: 651 649 5058
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Other ESTs: EST631492
Contact: Deborah A. Samac
Department of Plant Pathology
University of Minnesota
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 12)
Samac D.A., Dickman, M., Town, C.D., Van Aken, S., Utterback, T., Samac D.A., Dickman, M., Town, C.D., Van Aken, S., Utterback, T., Cheung, F. and Fraser, C.M.
ESTs from mycelia of Colletotrichum trifolii race 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes; Sordariomycetes incertae sedis; Phyllachorales; Phyllachoraceae; mitosporic Phyllachoraceae; Colletotrichum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: (gtA AtA CgA CtC ACt AtA ggg C).
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: debbys@puccini.crl.umn.edu
TIGR sequence name: MTSAA51TV More information
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Caenorhabditis elegans
                                        C51419.1 GI:2389176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: pBluescript SK+; Site 1: EcoRI; Site 2: /note="Vector: pBluescript SK+; Site 1: EcoRI; Site 2: EcoRI; isolate: 2sp2; cDNA was prepared from polyA+ enriched RNA The cDNA was ligated into Lambda gtil from Stratagene and packaged using Glgapack packaging extracts. An aliquot of the amplified library was used to transduce E. coli Y1090 and phage DNA was purified from a liquid lysate. The cDNA inserts were gel purified after EcoRI digestion and ligated into pBluescript SK+. Aliquots of the ligation were used to transform E. coli DH5alpha which
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/lab_host="DH5alpha"
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tissue_type="mycelia"
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Indels:
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Locus
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1 (bases 1 to 12)
Kohara,Y., Motohashi,T., Tabara,H., Watanabe,H., Sugimoto,A., Sano,M., Miyata,A. and Nishigaki,A.
Expression map of the C.elegans genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genome Biology Lab.
National Institute of Genetics
Yata 1111, Mishima, Shizuoka 411, Japan
Tel: 81-559-81-6854
Fax: 81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Caenorhabditis elegans
                                                                                                                                                                          Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
                                                                                                                                                                                                                                                  Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                            Oryza sativa
Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sativa cDNA clone 7LEAF--04-J19, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CF300273 12 bp mRNA linear EST 15-AUG-2003 7LEAF--04-J19.91 Rice leaf plasmid cDNA library II (7LEAF) Oryza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ykohara@lab.nig.ac.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis
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                                                                                                                                                                                                                                    Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CIGCGIATA 4
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                                                                                                                                    gin, Kyeonggi, Ko
: 82 31 330 6193
: 82 31 321 6355
                                                                                              bhnahm@ggbio.com, bhna
Location/Qualifiers
/organism="Oryza sativa"
/mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:4530"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Yuji Kohara unpublished cDNA"
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dev_stage="varied"
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                                                                                                                    bhnahm@bio.myongji.ac.kr
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CF311969
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KEYWORDS
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Genomics and Genetics Institute, Greendene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CF311969 12 bp mRNA linear EST 15-AUG-2003
ABF--07-H13.gl ABF3-overexpressing transgenic rice plasmid cDNA
library (ABF) Oryza sativa cDNA clone ABF--07-H13, mRNA sequence.
CF311969
CF311969.1 GI:33683730
                                                                                                                                                                                                                                                                                                                                                                                                                                         Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 12)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
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Oryza sativa
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                                                                                                                                                                                                                                                                                                                                                                                                                       bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
                                                                                                                     /note="Vector: pCR4-TOPO; Site 1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression line."
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/lab_host="E.coli DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid cDNA library (ABF)"
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/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
/mrh_cligoribonucleotides and then used as templates for
                                                                                                                                                                                                                                                                                                                                                    /organism="Oryza sativa"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                              tissue_type="leaf"
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/clone="ABF--07-H13"
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SOURCE

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CF329021 12 bp mRNA linear EST 18-AUG-2003 NACL-04-D03.g1 Rice callus plasmid cDNA library (NACL) Oryza sativa cDNA clone NACL--04-D03, mRNA sequence. CF329021
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABF--07-H13.gl ABF3-overexpressing transgenic rice plasmid cDNA library (ABF) Oryza sativa cDNA clone ABF--07-H13, mRNA sequence.

CF311969
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 12)
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Location/Qualifiers
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/mol type="mRNA"
/culTivar="Nackdong"
/db xref="taxon:4530"
/clone="ABF--07-H13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /dev_stage="14 days after germination"
/lab host="E.coli DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid
/clone lib="ABF3-overexpressing transgenic rice plasmid
/cDNA library (ABF)"
/note="Vector: pCR4-TOPO; Site 1: EcoRI; Leaf was dried
for 2hrs. Oligo-capped mRNA was reverse transcribed and
then used for PCR. mRNA was prepared from ABA-responsive
element binding transcription factor 3 overexpression
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., L.

Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                     Oryza sativa Oryza sativa Oryza sativa Embryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                        Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                           NACL--08-E07.gl Rice callus plasmid cDNA library (NACL) Oryza sativa cDNA clone NACL--08-E07, mRNA sequence. CF331951. GI:33812123
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)

Contact: Nahm B.H.
                                                                                                 Contact: Nahm B.H.
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                                       Yongin,
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, Kyeonggi, Ko
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tissue_type="callus"
/dev_stage="proliferated callus on 2N6 media for 30 days"
/lab_nost="E_coli DH10B"
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/mol_type="mRNA"
/cultivar="Nackdong"
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/clone="NACL--04-D03"
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                                                                                                                                                                                                                                                                                                                                                                                      Tel: (54-11) 4580/7255/7
Fax: (54-11) 4752-9639
Email: dBanchez@iib.unsam.edu.ar
                                                                                                                                                                                                                                                                                                                              On Jul 21, 2000 this sequence version replaced gi:6478057.
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Location/Qualifiers
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/cell_type="epimastigote"
/clone_lib="Trypanosoma cruzi random genomic library"
/note="Vector: pBS(-) (Stratagene); T. cruzi DNA was
randomly sheared using a nebulizer and the 1 to 2 Kb
                                                                                                /organism="Trypanosoma cruzi"
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/strain="CL-Brener"
                                                                   /clone="G35A7"
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/dev_stage="proliferated callus on 2N6 media for 30 days"
/lab_host="E.coli DH10B"
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/db_xref="taxon:4530"
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|mol_type="mRNA"
                                                                                 db_xref="taxon:5693"
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BH127723
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: Michele.morgante@usa.dupont.com
Sequences were trimmed to include only high quality bases; forward
and reverse reads were assembled when significant overlaps were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dupont Genomics
PO Box 6104, Newark, DE 19714-6104, USA
Tel: 302 631 2638
Fax: 302 631 2607
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Meyers,B.C., Tingey,S.V. and Morgante,M.
Abundance, distribution and transcriptional activity of repetitive elements in the maize genome
Genome Res. 11 (10), 1660-1676 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          clone G-1021 both, genomic survey sequence.
BH127723
BH127723.1 GI:14995555
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12 bp DNA linear GSS 23-JUL-2001
G-1021.r Maize Random Small-insert Genomic Library Zea mays genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Suite 200
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                                                                                                                                   /sex="hermaphrodite"
/tissue_type="leaf"
/cell_type="loaf"
/dev_stage="seedling"
/clone_lib="Maize Random Small-insert Genomic Library"
/note="Vector: pCR-Script; Total genomic DNA was
nebulized; ends were polished with Pfu polymerase and the
fragments cloned into pCR-Script."
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HincII site of the vector"
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'clone="G-1021"
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/strain="B73"
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RESULT 88
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PO Box 6104, Newark, I
Tel: 302 631 2638
Fax: 302 631 2607
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: Michele.morgante@usa.dupont.com
Sequences were trimmed to include only high quality bases;
and reverse reads were assembled when significant overlaps
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Suite 200
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Genome Res. 11 (10), 1660-1676 (2001)
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G-5a10.f Maize Random Small-insert Genomic Library Zea mays genomic
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Class: shotgun.
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Abundance, distribution and transcriptional activity of repetitive
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
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                                                                                                                                                                                                                                                                                                            /sex="hermaphrodite"
/tissue_type="leaf"
/cell_type="Young leaf"
/dev_stage="seedling"
/clone_lib="Maize Random Small-insert Genomic Library"
/note="Vector: pCR-Script; Total genomic DNA was
nebullzed; ends were polished with Pfu polymerase and i
fragments cloned into pCR-Script."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Zea mays"
/mol_type="genomic DNA"
/strain="B73"
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           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                EST; expressed sequence tag.
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                                                                                                                                                                                       AL043127;
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                                       Homo sapiens (human)
                                                                                         sapiens mRNA; EST DKFZp434D2223_r1
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Pooideae; Triticeae; Aegilops.
[Chases 1 to 12]
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CG677120
CG677120.1 GI:37506044
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tmf0875 tmf Aegilops tauschii
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Seq primer: T7
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Wheat Genetics Resource (
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/note="Vector: PCR 4Blunt-TOP; 0.8-1.2 kb methylation
filtered genomic DNA library"
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/dev_stage="shoot"
/lab_host="E. coli strain_DH5alpha"
/clone_lib="tmf"
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clone="tmf17C15"
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/mol_type="genomic DNA"
/strain="AL 8/78"
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S 66506-5502, USA
                                                                                           (from clone DKFZp434D2223)
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AA918967
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AA918667
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1 (bases 1 to 13)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP);
                                                                                                                                                                                          Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                        Homo sapiens (human)
Trace considered overall poor quality Insert Length: 1058 Std Error: 0.00
                                                           cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
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                                                 www-bio.llnl.gov/bbrp/image/image.html
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/dev_stage="adult"
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/mol_type="mRNA"
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DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
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                                                                                  Trace considered overall poor quality Seq primer: -40UP from Gibco High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 13)
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National Cancer Institute, Cancer Genome Ana
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AI744941.1 GI:5113229
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/mol_type="mRNA"
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                                                                                                                                                                                                                                            Email: sanjay_kumar-1@gsk.com
                                                                                                                                                                                                                                                                   Tel: 610-270-7245
Fax: 610-270-5598
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UW2109
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                                                                                                                                                                                                                           primer:
                 /tissue_type="cartilage"
/lab host="E.coli DH10 B"
/clone_lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPCRT I; Site_1: SalI; Si
Directional"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="tumor, 5 pooled (see description)"
/lab host="DH10B"
/clone_lib="NCI_CGAP_Ov23"
/clone_lib="NCI_CGAP_Ov23"
/note="Organ: ovary; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.35 kb. Tumor types include: mixed Mullerian tumor, papillary serous, clear cell, spindle cell. All are primary tumors, metastasis positive. Life Technologies catalog #: 11534-013"
                                                                                                                     /organism≈"Homo sapiens"
/mol_type="mRNA"
/db_xref≈"taxon:9606"
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Tetrahymena thermophila cDNA, mRNA secuence
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EST.
Tetrahymena thermophila
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Fax: 610-270-5598
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Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen
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HNC1-1-H7.R HNC (Human Normal Cartilage) Homo sapiens cDNA,
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/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
Directional"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="cartilage"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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Gowen, M. and
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Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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                                                                                                                                                                       University of Chicago
920 E. 58th Street, C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5009-0-1-H10.t.1 Chilcoat/Turkewitz CDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
                                                                                                                      Email: apturkew@midway.uchicago
                                                                                                                                           Tel: 773 702 4374 Fax: 773 702 3172
                                                                                                                                                                                                             Molecular Genetics and Cell Biology
                                                                                                                                                                                                                                    Contact: Turkewitz AP
                                                                                                                                                                                                                                                       Unpublished (2002)
                                                                                                                                                                                                                                                                      Frankel, J. and Klobutcher, L. Errain CU428.1, growing cells
                                                                                                                                                                                                                                                                                                        1 (bases 1 to 13)
Turkewitz, A.P., Ka
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University of Chicago
920 E. 58th Street, Chicago, IL 606:
Tel: 773 702 4374
Fax: 773 702 3172
Email: apturkew@midway.uchicago.edu
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             /organism="Tetrahymena
/mol_type="mRNA"
/strain="CU428.1"
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/db xref="taxon:5911"
/clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/clone_lib="Chilcoat SCH, Details on library
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beta vulgaris
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Trach
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicot
Caryophyllales; Amaranthaceae; Beta.
1 (bases 1 to 13)
Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,P.
Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
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13 bp mRNA linear EST 06-DEC-2002
E012394-024-013-F21-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone
024-013-F21 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADIS DNA core facility at MPIZ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: weisshaa@mpiz-koeln.mpg.de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Weisshaar B.
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primer: SP6; CATACGATTTAGGTGACACTATAG.
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Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database:http://gabi.rzpd.de"
                                                                                                                                                     /lab_host="EMDH10B"
/clone lib="MPIZ-ADIS-024-leaf"
/clone lib="MPIZ-ADIS-024-leaf"
/note="Vector: pcMVSPORT6; Site 1: Sall; Site 2: Notl;
/DNA library from sugar beet, library provided by KWS
CDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Binbeck, Germany, contact:
                                                                   SP6-Sall-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Sequencing granted in the context of the GABI-Beet
                                                                                                                                       b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="Chilcoat/Turkewitz_cDNA_(large_fraction)" /note="Vector: BlueScript2_SK+; Details on library preparation can be found in Chilcoat and Turkewitz_(2001) Proc. Natl. Acad. Sci_USA, 98: 8709-8713."
                                                                                                                                                                                                                                                                             /db_xref="GABI:186838"
/db_xref="taxon:161934"
/clone="024-013-F21"
/tissue_type="leaf"
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                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Beta vulgaris"
/mol tvoe="mRNA"
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ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Meg 10, 50829 Koeln, Germany
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Seq primer: SP6; CATACGATTTAGGTGACACTATAG.
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Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,
Drungowaki,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lel
and Radelof,U.
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllales; Amaranthaceae; Beta.
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E012680-024-020-D03-SP6 MPIZ-ADIS-024-storage root Beta vulgaris
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Insert Length: 13 Std Error: 0.00
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                   SP6-SalI-CCACGCGTCGG-5prime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de"
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/clone_lib="MPIZ-ADIS-024-storage root"
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CDNA_library_from_saltzucht_AG_Einbeck, Germany, contact
Kleinwanzlebener_Saltzucht_AG_Einbeck, Germany, contact
b.schulz@kws.de; cloning_sites_SalI-NotI, primer_sites_
b.schulz@kws.de; cloning_sites_SalI-NotI, primer_sites_b.
                                                                                                                                                                                                                            orientation:
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/db_xref="taxon:161934"
/clone="024-020-D03"
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Alignment Scores: Pred. No.:

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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
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1 (bases 1 to 13)
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BQ595423.1
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/note-Tvector: pcMVSPORT6; Site 1: SalI; Site 2: NotI;
cDNA library from sugar beet, library provided by KWS
CDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                        Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by
                                                                                                                                                                      RZPD/GABI-Primary database: http://gabi.rzpd.de"
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/lab_host="EMDH10B"
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/clone="024-022-N20"
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ol_type="mRNA"
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ot Beta vulgaris
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Insert Length: 13 Std Error: 0.00
Plate: 22 row: C column: 18
Seq primer: SP6; CATACGATTTAGGTGACACTATAG.
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Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,
Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Leb
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Max-Planck-Institute for Plant Breeding Research
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                                                                                                                                                                                                                                                                                             Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator:
                                                                                                                                                                                                                                                                                                                                            sp6-sall-ccacgcgrccg-sprime-cDNA-polyA-cc-NotI-T7;
                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="MPIZ-ADIS-024-developing root"
/note="Vector: pCMVSPORT6; Site 1: Sal1; Site 2: Not1;
/DNA library from sugar beet, library provided by KWS
Kleinwanzlebener Sastrucht AG Binbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                           Prof. Christian Jung; Sequo
RZPD/GABI-Primary database:
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db_xref="taxon:161934"
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|mol_type="mRNA"
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3.00
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                                                                                                       Mismatches:
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CF299938/c
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EST.
Theobroma cacao (cacao)
Oryza sativa
Oryza sativa
                            EST
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Jones, P.G., Allaway, D., Gilmour, D.M., Harris, C., Rankin, D.
Retzel, E.R. and Jones, C.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Malvales; Malvaceae; Byttnerioideae;
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13 bp mRNA linear EST 05-DE Cac BL 1304 Cac BL (Bean and Leaf from Amelonardo type Cacao)
TheoDroma cacao cDNA clone Cac BL 1304 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3d Dundee Road, Slough, Berkshire,
Tel: +44 1664 416644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Jones, Paul
Masterfoods
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Planta 216 (2), 255-264 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                     /clone lib="Cac_BL (Bean and Leaf from Amelonardo type Cacao)"
                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: pBK-CMV; Bean and leaf tissue from an Amelonado type Cacao tree."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         db_xref="taxon:3641"
/clone="Cac_BL_1304"
/tissue_type="Mature leaf and mature bean"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism≈"Theobroma cacao"
/mol_type="mRNA"
/strain="Amelonado type"
                                                                                                                                                                                                                                                                                                                                                                                                                                 dev_stage="maturity"
lab_host="XL-1 Blue MRF'"
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13 bp mRNA linear EST 15-AUG-2003
HDA1--02-L18.g1 OsHDAC1-overexpressing transgenic rice lambda phage
cDNA library I (HDA1) Oryza sativa cDNA clone HDA1--02-L18, mRNA
                                                                                                                           Contact: Nahm B.H.

Genomics and Genetics Institute, Greendene Biotech I of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                   1 (bases 1 to 13)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,

Song,S.I., Xim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                   Oryza sativa
Oryza sativa
                                                                 Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence.
CF306112
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Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
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    Location/Qualifiers
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Fax: 82 31 321 6355
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Unpublished (2003)
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bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
1. .13
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/db xref="taxon:4530"
/db xref="taxon:4530"
/clone="TLEAF-04-C12"
/tissue_type="leaf"
/dev_stage="7 days after germination"
/dev_stage="7 days after germination"
/lab_host="E.coli DH10B"
/clone_lib="Rice_leaf_plasmid_cDNA_library_II_(7LEAF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA_was_capped_with_oligoribonucleotides_and_then_used_as_templates_for_RT-PCR."
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|mol_type="mRNA"
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                                                                                                                                                            Biotech Inc.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H. and Radelof,U.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CF543088 13 bp mRNA linear EST 22-SEP-2003 S014680-024-030-P08-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone 024-030-P08 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                             ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
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                                                                                                                                                                                                                                                                                                                                                                                     Contact: Weisshaar B
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                                                                                                                                                                                                                        primer: SP6
                                                                                                                                                                                                                                           /note-"Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: /note-"Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: XhoI; Callus was treated with ABA(20um) for 1hour. cDNA was inserted into lambda Uni-ZAP XR vector at 5' end with EcoRI and 3' end with XhoI site. mRNA was derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /dev stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="E.coli SOLR"
/clone_lib="OsHDAC1-overexpressing transgenic rice lambda
phage cDNA library I (HDA1)"
                                                                                                                                          organism="Beta vulgaris"
/mol_type="mRNA"
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                    clone="024-030-p08"/tissue_type="leaf"
                                                            /db_xref="GABI:936675"
/db_xref="taxon:161934"
                                                                                                                       cultivar="KWS2320 (double
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/clone="HDA1--02-L18"
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/mol_type="mRNA"
/cultivar="Nackdong"
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Alignment ORIGIN

Мо.:

RESULT 103

PUBMED COMMENT

JOURNAL MEDIINE

TITLE

FEATURES

100.00% 2.73e+06 3.00

Matches: Conservative: Mismatches:

00 4 13

REFERENCE AUTHORS

SOURCE

ORGANISM

KEYWORDS ACCESSION DEFINITION CF543088/c

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Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database:http://gabi.rzpd.de"
                                                                                                                                                                                                                                                                                                                             /clone_lib="MPIZ-ADIS-024-leaf"
/note="Vector: pCMVSPORT6; Site_1: Sall; Site_2: NotI;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact
                                                                                                                                         SP6-Sall-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Sequencing granted in the context of the GABI-Beet
                                                                                                                                                                                                                                                                                                                                       contact:
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US-09-966-880A-8 (1-198) x CF543088 (1-13)
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Glycine max (soybean)
Glycine max
Glycine max
Viridiplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        171 ArgLeuSer 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12 CGCCTGTCA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gmrhRww3-07_G12_1_084 Soybean root hair s
gmrhRww3 Glycine max cDNA, mRNA sequence.
CF921303.1 GI:38192097
                                                                                                                                                                                                                                                                                                                                                                 Contact: Gary Stacey
University of Missouri
108 Waters Hall, Columbia, MO 65211, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Glycine.

1 (bases 1 to 13)
Scheffler, B.B., Huang, S., Liu
Scheffler, B.B., Huang, S., Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
                                                                                                                                                                                                                                                            Single pass sequence
Seq primer: T7.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Expressed sequence tags
                                                                                                                                                                                                                                                                                                           Email: staceyg@missouri.edu
                                                                                                                                                                                                                                                                                                                                Fax: 573-882-0588
                                                                                                                                                                                                                                                                                                                                                  Tel: 573-884-4752
/note="Organ: root hairs; Vector: pCR2-1 Topo; cDNA clones generated from soybean root hair tissue treated with Bradyrhizobium japonicum for 3 hours."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                       /tissue_type="root hairs"
/clone_Tib="Soybean root hair subtracted cDNA library
                                                                                                                                 /organism="Glycine max"
/mol_type="mRNA"
/cultivar="Williams 82"
/db_xref="taxon:3847"
                                                                                                                                                                                                                                        Socation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.00%
100.00%
1.52%
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root hair subtracted cDNA library
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RESULT 106
HSM003815/c
ID HSM0038:
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AUTHORS
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DB:
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                                                                                                                                                                                                                              Query Match:
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                                                                                                                                                                                                                                                                   Percent Similarity:
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                   HSM003815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   source
                                                                                                                                                                                                                                                                                                          No.:
                                                                                                                             129 LeuHisArg 131
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A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Tex: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: TDNA tagged.
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Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SerPheSer 43
                                                                                              CTTCACCGC
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                   standard; mRNA; EST; 14 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0808
_003378 Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Ārabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fargment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'mol_type="genomic DI
'strain="Columbia 0"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         db_xref="taxon:3702"
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100.00%
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Matches:
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                                                                                                                                                                                                                                                               Query Match:
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Best Local Similarity:
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No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No sl sequence available
This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerwe
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                                                                                                                                                                                                                                                                          Sequence 14 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clone from S. Wiemann, sequenced by Qiagen within the cDNA sequencing consortium of the German Genome Project
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Eutheria; Primates; Catarrhini;
Homo sapiens (human)
                                                                12-MAR-1999
12-MAR-1999
                                                                                                AL039902.1
                                                                                                                     AL039902;
                                                                                                                                           HSM004378
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Duesterhoeft A., Lauber J., Mewes W., Gassenhuber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens mRNA; EST DKFZp434F1810_r1 (from clone DKFZp434F1810)
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12-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expressed sequence tag.
                                          sapiens mRNA;
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                                                                                                                                                                                                          PheArgThr 195
                                                                                                                                                                                    TTCCGGACC
                                                                                                                                         standard; mRNA; EST; 14
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                                                                                                                                                                                                                                                                                                                                                                        /orgānism="Homo sapiens"
/clone="DKF2p434F1810"
/clone_lib="434 (synonym: h
DH10B; sites NotI + SalI"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                          3 A; 4 C; 4 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                              'tissue_type="testis"
                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:9606"
/mol_type="mRNA"
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,
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59, Last updated, Version
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100.00%
100.00%
1.52%
                                         EST DKFZp434G2412_r1
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                                                                                                                                                                                                                                                                                                                                                                                                                               _type="mRNA"
                                                               Created)
Last upd
                                                               updated,
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                                                                                                                                                                                                                               (1-14)
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Matches:
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                                                                                                                                                                                                                                                               Mismatches:
Indels:
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                                                                                                                                           ₽₽.
                                                               Version 1)
                                          (from clone DKFZp434G2412)
                                                                                                                                                                                                                                                                                                                                                                                              htes3). Vector pSport1; host
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clone is available at the RZPD in Berlin
se contact the RZPD: Ressourcenzentrum, Ho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       193 PheArgThr 195
GlaxoSmithKline
709 Swedeland Road,
Tel: 610-270-7245
Fax: 610-270-5598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A
                                                                                                                                                                                                                                                                                                                                                                                                                                                             10
                                                                                                                      sequenced tags (ESTs) each from adult he osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (200)
                                                                                                                                                                                                   Kûmar,S., Connor,J.R., Dodds,R.A., Halsey,W., Van Horn,M., Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen.
                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 14)
                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                               HNC27-1-D6.R
                                                                          Contact: Sanjay Kumar
                                                                                                                                                       Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and
                                                                                                                                                                                                                                                                                                                                 BG924475.1 GI:14318998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 S. Wiemann, sequenced by Qiagen within the cDNA consortium of the German Genome Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/clone="hKFZp434G2412"
/clone_lib="434 (synonym: htes3). Vector pSport1;
bH10B; sites NotI + SalI"
/dev_stage="adult"
/tissue_type="testis"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:9606"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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5009-0-55-C12.t.2 Chilcoat/Turkewitz cDNA (1
Tetrahymena thermophila cDNA, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                              Tel: 773 702 4374 Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                       Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 606
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                                                                                                                                                                                                                                                                                                                              Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Turkewitz AP
                                                                                                                                                                                                                                                                                                                                                                                                                                        EST from Tetrahymena thermophila, strain CU428.1, Unpublished (2002)
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Turkewitz, A.P., Karrer, K.M., Jahn, C.,
Frankel, J. and Klobutcher, L.
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/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/lab_host="B.coli DH10 B"
/clone_lib="HNC (Human Normal Cartilage)"
/note="Vector: psport I; Site_1: SalI; Site_tional"
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1 (bases 1 to 14)
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/note="Vector: pCMVSPORT6; Site 1: Salte 2: Not1;
/DATA library from sugar beet, library provided by KWS
CDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SP6-Sall-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           orientation:
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/mol_type="mRNA"
/cultivar="KWS2320 (doubl
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/Lab_host="EMDH10B"
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
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Contact: Weisshaar B
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1 (bases 1 to 14)

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/note="Vector: pCMVSPORT6; Site 1: SAI:, Site 2: Not1;
/note="Vector: pCMVSPORT6; Site 1: SAI:, Site 2: Not1;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SP6-SalI-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet
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/db_xref="taxon:161934"
/clone="024-019-A07"
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/mol_type="mRNA"
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'lab_host="EMDH10B"
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Tracheophyta;

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14 bp mRNA linear EST 06-DEC-2002 S013408-024-026-P02-SP6 MPIZ-ADIS-024-developing root Beta vulgaris cDNA clone 024-026-P02 5-PRIME, mRNA sequence.
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllales; Amaranthaceae; Beta.
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EST.
             BQ593808 14 bp mRNA linear EST 06-DEC-2002 E012763-024-026-007-SP6 MPIZ-ADIS-024-developing root Beta vulgaris CDNA clone 024-026-007 5-PRIME, mRNA sequence.
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/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Notl;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Notl;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b:schulz@kws.de; cloning sites Sall-Notl, primer sites and
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/lab_host="EMDH10B"
/clone_lib="MPIZ-ADIS-024-deve
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'clone="024-026-P02"
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CA798290

CAC_BL_611 Cac_BL (Bean and Leaf from Amelonardo type Cacao)
Theobroma cacao cDNA clone Cac_BL_611 5', mRNA sequence.
CA798290
CA798290.1 GI:26055376
EST
Theobroma cacao (cacao)
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SP6-SalI-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Note:
Sequencing granted in the context of the GABI-Beet
Sequencing granted in the context of the GABI-Beet
project, local PI: Dr. Katharina Schneider, coordinator:
Prof. Christian Jung; Sequence submission managed by
RZPD/GABI-Primary database: http://gabi.rzpd.de"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="MPIZ-ADIS-024-developing root"
/note="Vector: pCMVSPORT5; Site_1: Sall; Site_2: Not1;
cDNA library from sugar beet, library provided by KWS
cDNA library from sugar beet, library provided by contact
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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/lab_host="EMDH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="GABI:193044"
/db_xref="taxon:161934"
/clone="024-026-007"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Beta vulgaris"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cultivar="KWS2320 (double haploid, monogerm breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              schulz@kws.de; cloning sites SalI-NotI, primer sites and
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EST 05-DEC-2002

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CA853334/c
                                                        JOURNAL COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Masterfoods
3d Dundee Road, Slough, Berkshire,
Tel: +44 1664 416644
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Jones, P.G., Allaway, D., Gilmour, D.M., Harris, C., Rankin, D.
Retzel, E.R. and Jones, C.A.
                                                                    1 (bases 1 to 14)
Alkharouf, N.W., Khan, R. and Matthews, B.F.
Analysis of expressed sequence tags from roots of resistant soybean infected by the soybean cyst nematode
Unpublished (2002)
                                                                                                                                                                                                                                       CA853334.1 GI:33390127
EST.
Glycine max (soybean)
Glycine max
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Soybean Genomics and Improvement Laboratory (SGIL) US Department of Agriculture (USDA), ARS, PSI Bldg.006, Rm 118, 10300 Baltimore Ave., Beltsville
                                                       Contact: Alkharouf, N.W.
                                                                                                                                                                                  Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: pBK-CMV; Bean and leaf tissue from an
Amelonado type Cacao tree."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="Mature leaf and mature bean"
/cell_type="Whole organ"
/dev_stage="maturity"
/lab_host="XL-1 Blue MRF'"
/clone_lib="Cac_BL (Bean and Leaf from Amelonardo type
Cacao)"
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'strain="Amelonado type"
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CF278327/c
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KEYWORDS
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14ETL--04-D06.b1 Rice etiolated leaf plasmid cDNA library (14ETL)
Oryza sativa cDNA clone 14ETL--04-D06, mXNA sequence.
CF278327
                                                                                                                                                                                                                                                                                                     Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 14)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Losong,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)

Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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Oryza sativa
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Fax: 301 504 5728
                                                                                                                                                                                                                                                                                                                                                             Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CF278327.1
                                                                                                                                                                                                                                                               bhnahm@ggbio.com, bhna
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="Roots"
/dev_stage="Seedlings"
/clone_lib="CDNA_Peking_library_12hr_SCN3"
/note="Vector: pBluescript_SK-; cDNA_clones_from mRNA_extracted_from_roots_of_soybean_cv_Peking_12_hrs_after_infection_by_SCN_race_3. These_are_cloned_in_pBluescript_SK-_phagemid."
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/clone="B07A06"
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/cultivar="Peking"
with oligoribonucleotides
                                                                                                                                                                                /organism="Oryza sativa"
/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Glycine max"
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            note="Vector: pCR4-TOPO; Site 1: EcoRI; mRNA was capped with oligoribonucleotides and then used as templates for
                                                      clone lib="Rice etiolated leaf plasmid
                                                                                    tissue_type="leaf"
dev_stage="14 days after germination"
lab_host="E.coli DH10B"
                                                                                                                                            db_xref="taxon:4530"
clone="14ETL--04-D06"
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RESULT 118
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Oryza sativa
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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7LEAF--05-B01.gl Rice leaf plasmid cDNA library II (7LEAF) Oryza
sativa cDNA clone 7LEAF--05-B01, mRNA sequence.
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Location/Qualifiers
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1 (bases 1 to 14)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)

Contact: Nahm B.H.
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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HDA1--05-D06.gl OsHDAC1-overexpressing transgenic rice lambda phage
CDNA library I (HDA1) Oryza sativa cDNA clone HDA1--05-D06, mRNA
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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7LEAF--05-B01.gl Rice leaf plasmid cDNA library II (7LEAF) Oryza
Batiya CDNA clone 7LEAF--05-B01, mRNA sequence.
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Location/Qualifiers
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with oligoribonuclectides and then used as templates for
RT-PCR."
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Fax: 82 31 321 6355
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
Contact: Nahm B.H.
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
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82 31 330 6193
82 31 321 6355
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Location/Qualifiers
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HDA1--06-N23.gl OsHDAC1-overexpressing transgenic rice lambda phage
cDNA library I (HDA1) Oryza sativa cDNA clone HDA1--06-N23, mRNA
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: 82 31 330 6193
: 82 31 321 6355
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Location/Qualifiers
                  /clone_lib="OsHDAC1-overexpressing transgenic rice lambda phage cDNA library I (HDA1)"
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rice Histone Deacetylase overexpression line."
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CF328966.1 GI:33806172
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
Contact: Nahm B.H.
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82 31 330 6193
82 31 321 6355
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone lib="Rice callus plasmid cDNA library (NACL)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with oligoribonuclectides and then used as templates for
RT-PCR."
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/mol_type="mRNA"
/cultivar="Nackdong"
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[lab_host="B_coli DH10B"]
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Bukaryota; Viridiplantae; Streptophyta; core endicots;
Spermatophyta; Magnoliophyta; endicotyledons; core endicots;
rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
Contact: Nahm B.H.
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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Oryza sativa
                              Tel: 573-884-4752
Fax: 573-882-0588
                                                                                 Unpublished (2003)
Contact: Gary Stacey
University of Missouri
108 Waters Hall, Columbia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Yongin, Kyeonggi, Korea
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Scheffler, B.E., Huang, S.,
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Location/Qualifiers
  staceyg@missouri.edu
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/dev_stage="proliferated callus on 2N6 media for 30 days"
/dev_stage="proliferated callus on 2N6 media for 30 days"
/lab_host="E.coli DHIOB"
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/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA_was_capped
/note="library_collision: pcR4-TOPO; site_1: EcoRI; mrNA_was_
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/cultivar="Nackdong"
/db_xref="taxon:4530"
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                                                                                                                                                                                                                                                                                                                                                 Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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Seq pri
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Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L.,
Shinn,P., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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14 bp DNA linear GSS 03-OCT-200 SALK_00178B Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_001788, genomic survey sequence.
BH169716
                                                                                                                                                                                                                                                                  Class: TDNA tagged
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GSS.
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                                                                                                                                                                                                                                                                                                            is single pass sequence recovered
                                                                                                                                                                                                                                                                                                                                 ecker@salk.edu
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/clone_lib="Ārabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used or
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/clone="SALK_001788"
                                                                                                                                               /strain="Columbia 0"
'db xref="+---
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/clone_lib="Soybean root hair subtracted cDNA library
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VERSION
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Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
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Tel: 858 453 4100 x1752
Fax: 858 558 6379
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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SALK 001788 Arabidopsis thaliana TDNA insertion lines Arabidopsis
thaliana genomic clone SALK_001788, genomic survey sequence.
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/clone lib="Arabidopsis thaliana TDNA insertion lines"
/clone lib="Arabidopsis thaliana lines
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
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/strain="Columbia 0"
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                                                                                                                                                       Sequence 15
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                                                                                                                                                                                                                                                                         sequencing consortium of the s1 sequence also available
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                                                                                                                                                                                                                                                                                                    MIPS,
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone_lib="434 (Bynonym: htes3).
DH10B; sites Not! + Sal!"
/dev_stage="adult"
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/organism="Homo sapiens"
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sequencing consortium of
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Please contact the RZPD: Ressourcenzentrum, H
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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Homo sapiens (human)
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/clone="Dxr2p3420823"
/clone_lib="434 (synonym: k
DH10B; sites NotI + SalI"
/dev_stage="adult"
/figue_type="adult"
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Indels:
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e@rzpd.de
                                                       clone DKFZp434L1223)
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Ξ Eutheria;

Eukaryota; Metazoa;

; Metazoa; Chordata; Craniata; Vertebrata; Primates; Catarrhini; Hominidae; Homo.

Euteleostomi; Mammalia;

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Best Local Similarity:
Query Match:
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This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, Heubnerweg Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                Clone from S. Wiemann, sequenced by sequencing consortium of the German No 81 sequence available
                                                                                MIPS,
                                                                                                                                                                                                                                                                                                                                 HSM008148
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Butheria; Primates; Catarrhini; Hominidae; Homo.
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Please contact the RZPD: Ressourcenzentrum, H
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                                                                                                                  BLum
                                                                                                                                                                                Homo sapiens (human)
                                                                                                                                                                                                        EST; expressed sequence
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                                                                               Klopferspitz
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DH10B; sites NotI + SalI"
/dev_stage="adult"
/tissue_type="testis"
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ERMANY; Email: clone@rzpd.de
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D-82152 Martinsried, GERMANY
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JOURNAL
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Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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AL931094 NAP1 Anopheles gambiae
seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: +49 6221 387-440 Fax: +49 6221 387-306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     European Molecular Biology Laboratory Meyerhofstrasse 1, 69117 Heidelberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact:
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Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Christophides, G.K.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       christop@embl-heidelberg.
P72 row: D column: 06.
Location/Qualifiers
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/clone="DKF2p434M2423"
/clone_lb="434 (synonym: k
/dl0B; sites NotI + SalI"
/dev_stage="adult"
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                                                  library constructed from a mixture of Anopheles gambiae developmental stages according to: Bonaldo, Lennon & Soares (1996): Normalization and Subtraction: Two
                                                                                                                                     /Clone="NAP1-P72-D-06-5"
/lab_host="E. coli DH10B"
/clone lib="NAP1"
/note="Vector: pT7T3D-Pac (Pharmacia); Site_1: NotI;
/notI;
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Approaches To Facilitate Gene Discovery, 6, 791-806."
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db_xref="taxon:7165"
                                                                                                                                                                                                                                                                                                                                                                                                       organism="Anopheles
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AW059513 15 bp mRNA linear EST 23-AUG-2000 HuTH.bsst.dnc15.final.cluster_2_(36) DNC15 Homo sapiens cDNA similar to ribosomal protein $17, mRNA sequence.
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Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida;
Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /sex="hermaphrodite, male"
/tissue_type="whole animal"
/dev_stage="varied"
/clone_lib="Yuji Kohara unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="mRNA"
/strain="CB1489 him-8(e1489)"
/db_xref="taxon:6239"
/clone="yk546e2"
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Kohara unpublished cDNA Caenorhabditis elegans cDNA
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
                                                                         Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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LYNX Therapeutics, Inc.
LYNX Therapeutics, Inc.
25861 Industrial Blvd., Hayward, CA 94545, USA
Tel: 510 670 9338
Fax: 510 670 9302
Fax: 510 670 9302
Sequence obtained from LYNX Therapeutics Megasort technology.
Collected from the down-regulated gate. Consensus sequence of 36
sequences in cluster.
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Brenner, S., Williams, S.R., Vermass, E.H., Storck, T., Moon, K., McCollum, C., Mao, J.I., Kirchner, J.J., Eletr, S., DuBridge, R.B., Burcham, T. and Albrecht, G.
Burcham, T. and Albrecht, G.
In vitro cloning of complex mixtures of DNA on microbeads: Physical separation of differentially expressed cDNAs separation of differentially expressed cDNAs Proc. Natl. Acad. Sci. U.S.A. 97 (4), 1665-1670 (2000)
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2819953.3prime NIH_MGC_7
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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/note="Vector: pCR2.1; Cloning of PCR products from micro-beads carrying 3' end of down-regulated cDNA.cells non-induced (treated with DMSO only). "
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/mol_type="maXN,"
/mol_type="maXN,"
/db xref="rtaxon:9606"
/cell_type="monocyctic leukemia"
/cell_line="THP-1 (TIB-202)"
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Identification and initial characterization of 5000 expressed sequenced tags (RSTs) each from adult human normal and osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001)
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                                                                                                                               Lark, M.W
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Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen,M.
                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone lib="NIH MGC 7"
/clone lib="NIH MGC 7"
/clone lib="NIH MGC 7"
/note="Organ: lung; Vector: pOTB7; Site 1: XhoI; Site 2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5; adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
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/tissue_type="small cell carcinoma"
/cell_line="MGC3"
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/mol_type="mRNA"
/ac__
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Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen,M. and
Lark,M.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 15)
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15 bp mRNA linear EST 06-NOV-2
HNC5-1-B1.R HNC (Human Normal Cartilage) Homo sapiens cDNA, mRNA
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Fax: 610-270-5598
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Seq primer: T7.
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UW2109
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Seq primer: T7.
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/clone_lib="HOA (Human Osteoarthritic Cartilage)"
/note="Vector: pSPORT I; Site_1: Sall; Site_2: Not Directional"
/tissue_type="cartilage"
/lab_host="3.coli DH10 B"
/clone_lib="HNC (Human Normal Cartilage)"
                                                       organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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mol_type="mRNA"

db_xref="taxon:9606"
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Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen,M. and
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imer: T7.
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/lab_host="E.coli DH10 B"
/clone_lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
Directional"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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US-09-966-880A-8 (1-198) x BM396203
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Turkewitz,A.P., Karrer,K.M., Jahn,
Frankel,J. and Klobutcher,L.
EST from Tetrahymena thermophila,
Unpublished (2002)
                                                                                                                                                                                               Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota, Alveolata, Ciliophora, Oligohymenophorea,
Hymenostomatida, Tetrahymenina, Tetrahymena.
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Tel: 773 702 4374
Fax: 773 702 3172
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Turkewitz, A.P., Karrer, K.M., Jahn, C.,
Frankel, J. and Klobutcher, L.
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Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 60637, USA
                                                                                                                                                                                                                                                                                                                                                    5009-0-18-G08.t.1 Chilcoat/Turkewitz cDNA (large Tetrahymena thermophila cDNA, mRNA sequence.
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Seq primer: T3.
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                                                                  Contact: Turkewitz AP
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University of Chicago
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/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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strain="CU428.1"
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                                                                                                               strain CU428.1,
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RESULT 140
BM398486
                Percent Similarity:
Best Local Similarity:
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DB:
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JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota, Alveolata, Ciliophora, Oligohymenophorea,
Hymenostomatida, Tetrahymenina, Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1009-0-45-H01.t.2 Chilcoat/Turkewitz cDNA (large fraction)
Tetrahymena thermophila cDNA, mRNA sequence.
BM398486
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                                                                                                                                                                                                                                                                                                   University of Chicago
920 E. 58th Street, Chicago,
Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                              Molecular Genetics and Cell Biology
                                                                                                                                                                                                                                                                                                                                                                            Contact: Turkewitz AP
                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 15)
Turkewitz, A.P., Karrer, K.M.,
Frankel, J. and Klobutcher, L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BM398486.1 GI:18198539
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                                                                                                                                                                                                                                                                                                                                                                                                        from Tetrahymena thermophila, strain CU428.1, growing cells
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773 702 3172
                                                                                                                                                                                                                                                                                      apturkew@midway.uchicago.
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                                                                                                                   /db_xref="taxon:5911"
/clome_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/clome_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/clome_lib="Chilcoat/Turkewitz cDNA (large fraction)"
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preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                                                                                                                             /mol_type="mRNA"
/strain="CU428.1"
                                                                                                                                                                                                                         organism="Tetrahymena thermophila"
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/strain="CU428.1"
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Length:
Matches:
Conservative:
Mismatches:
Indels:
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AUTHORS
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BM415446
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Best Local Similarity:
Query Match:
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EST.
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                                                                                                     BQ584986
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Fax: 919.515.9500
                                                          Beta vulgaris
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US-09-966-880A-8 (1-198) x BM415446 (1-15)
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Beta vulgaris
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllales; Amaranthaceae; Beta.
                                                                                                                                                                                                                                                   BQ584986 15 bp mRNA linear E011826-024-002-K24-SP6 MPIZ-ADIS-024-inflorescence CDNA clone 024-002-K24 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: warthog@unity.ncsu.edu
GT11-4PCN F_D05_PCN_4_F_042.ab1.
Location/Qualifiers
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Mixed Stage EST's from Globodera pallida,
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Globodera pallida
Eukaryota; Metazoa; Nematoda; Chromadorea; Tylenchida; Tylenchina;
Tylenchoidea; Heteroderidae; Heteroderinae; Globodera.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BM415446.1 GI:18382117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="mRNA"
/db_xref="taxon:36090"
/clone_ib=-Mixed_Stage_EST's from Globodera_pallida,
potato_cyst_nematode"
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                                                                                                                                                                                                    GI:26114563
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lida, the potato cyst r
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LOCUS
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ORGANISM
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ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 15)
Herwig, S., Schulz, B., Weisshaar, B., Hennig, S., Steinfath, Drunggowski, M., Stahl, D., Wruck, W., Menze, A., O'Brien, J., and Radelof, U.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
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Insert Length: 15 Std Error: 0.00
Plate: 2 row: K column: 24
Seq primer: SP6; CATACGATTTAGGTGACACTATAG.
Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach, and Radelof,U.
                                                              Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Caryophyllales; Amaranthaceae; Beta.

1 (bases 1 to 15)
                                                                                                                                                   Beta vulgaris
Beta vulgaris
                                                                                                                                                                                                                                                            BQ588286 15 bp mRNA linear EST 06-DEC-2002
E012308-024-008-J22-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone
024-008-J22 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                      BQ588286.1 GI:26117869
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="MPIZ-ADIS-024-inflorescence"
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not1;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not1;
CDNA library from sugar beet, library provided by KWS
Cleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-NotI, primer sites and
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/lab_host="EMDH10B"
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/db_xref="taxon:161934"
/clone="024-002-K24"
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Matches:
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core ev
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Contact: Weisshaar
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    Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes plant J. 32 (5), 845-857 (2002) 22362189
                                                                                                           Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Leh
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Seq primer: SP6; CATACGATTTAGGTGACACTATAG
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Max-Planck-Institute for Plant Breeding Research
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1 (bases 1 to 15)
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Insert Length: 15 Std Error: 0.00
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/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not1;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not1;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Binbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-NotI, primer sites and
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/mol_type="mRNA"
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eudicots;
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1 (bases 1 to 15)

Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Dringowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
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BQ591870.1 GI:26121453
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15 bp mRNA linear EST 06-DEC-2002
E012581-024-016-M20-SP6 MPIZ-ADIS-024-storage root Beta vulgaris
CDNA clone 024-016-M20 5-PRIME, mRNA sequence.
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Plate: 15 row: K column: 22
Seq primer: SP6; CATACGATTTAGGTGACACTATAG.
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Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant
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/note="Vector: pcMVSPORT6; Site_3: Not1;
/note="Vector: pcMVSPORT6; Site_3
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|mol_type="mRNA"
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'db_xref="taxon:161934"
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BQ595631.1 GI:26125214
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Plate: 16 row: M column: 20
Seq primer: SP6; CATACGATTTAGGTGACACTATAG
                  Contact: Weisshaar B
ADIS DNA core facility at MPIZ
MMAX-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
                                                                                                                                                        Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet great J. 32 (5), 845-857 (2002)
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Fax: 00492215062851
Email: weisshaa@mpiz-koeln.mpg.de
Fax: 00492215062851
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SP6-SalI-CCACGCTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note
Sequencing granted in the context of the GABI-Beet
Sequencing granted in Katharina Schneider, coordinator:
project, local PI: Dr. Katharina Schneider, coordinator:
prof. Christian Jung; Sequence submission managed by
RZPD/GABI-Primary database: http://gabi.rzpd.de"
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Kleinwanzlebener_Saatzucht_AG_Binbeck, Germany, contact:
b.schulz@kws.de; cloning_sites_Sall-NotI, primer_sites_and
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weisshaa@mpiz-koeln.mpg.de Length: 15 Std Error: 0.00

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REFERENCE
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CA796369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Theobroma cacao

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Eukaryota; Varidiplantae; Streptophyta; Embryophyta; Tracheophyta;

Energy to the control of the cont
                                                                                                                                                                                           3d Dundee Road, Slough, Berkshire, Tel: +44 1664 416644
Email: Paul.Jones@eu.effem.com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EST.
Theobroma cacao (cacao)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CA796369 15 bp mRN/Cac_BL_3383 Cac_BL (Bean and Leaf from Theobroma cacao cDNA clone Cac_BL_3383
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Seq primer:
                                                                                                                                                                                                                                                                                                                         Contact: Jones, Paul
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                                                                                                                                                                                                                                                                                                                                                                                                                     Planta 216 (2), 255-264 (2002)
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                                                                                                                                                          il: Paul.Jones@eu.effem.com
primer: T3.
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/note="Vector: pCMVSPRT6; Site 1: SAI; Site 2: Not!;
/note="Vector: pCMVSPRT6; Site 1: SAI; Site 2: Not!;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sp6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de"
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r: SP6; CATACGATTTAGGTGACACTATAG.
Location/Qualifiers
/organism="Theobroma cacao"
/mol_type="mRNA"
/strain="Amelonado type"
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'clone="024-022-B04"
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|mol_type="mRNA"
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lab_host="EMDH108"
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      Alignment Scores:
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L.

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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ABF1--03-K24.g1 ABF3-overexpressing transgenic rice lambda CDNA library (ABF1) Oryza sativa CDNA clone ABF1--03-K24,
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Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Oryza sativa"
mol type="mRNA"
/cultivar="Nackdong"
/db xref="taxon:4530"
/clone="ABP1--03-K24"
/tissue_type="leaf"
/dev_stage="14 days after germination"
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/lab_host="S.coli SOLR"
/clone_lib="ABF3-overexpressing transgenic rice lambda
/clone_lib="ABF3-overexpressing transgenic rice lambda
phage_CDNA library (ABF1)"
/note="Vector: pBluescript_SK(+); Site_1: EcoRI; Site_2:
/note="Vector: pBluescript_SK(+); Site_1: EcoRI; Site_2: EcoRI; S
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/clone="Cac_BL_3383"
/tissue_type="Mature leaf and mature bean"
/cell_type="Whole organ"
/dev_stage="maturity"
/lab_host="XL-1 Blue MRF'"
/clone_lib="Cac_BL (Bean and Leaf from Amelonardo type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Vector: pBK-CMV; Bean and leaf tissue from Amelonado type Cacao tree."
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AUTHORS
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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                                                                      ProCysTyr 88
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                                                                                                                                                                                                                                                              /note="Vector: DCR4-TOPO; Site 1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression line."
                                                                                                                                                                                                                                                                                                                                      /clone lib="OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD)"
                                                                                                                                                                                                                                                                                                                                                                   /dev_stage="proliferated callus on 2N6 media for 2 weeks"
lab_host="E.coli_DH10B"
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/mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:4530"
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Oryza sativa
Oryza sativa
Oryza sativa
Coryza sativa
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 15)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
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HD--07-L05.g1 OsHDAC1-overexpressing transgenic rice plasmid cDNA
Library (HD) Oryza sativa cDNA clone HD--07-L05, mRNA sequence.
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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CF317855
CF317855.1 GI:33689616
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com,
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dev_stage="proliferated callus on 2N6 media for 2 weeks"
lab_host="E_coli_pH10B"
/clone_lib="0sHDAC1-overexpressing transgenic rice_plasmid
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/mol_type="mRNA"
/cultivar="Nackdong"
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                                                                                               Contact: Nahm B.H.

Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University

Yongin, Kyeenggi, Korea

Tel: 82 31 330 6193

Fax: 82 31 321 6355

Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CF324208 15 bp mRNA linear EST 18-AUG-2003
HDN--05-018.gl OsHDACl-overexpressing transgenic rice lambda phage
cDNA library II (HDN) Oryza sativa cDNA clone HDN--05-018, mRNA
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82 31 321 6355
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/organism="Oryza sativa"
/mol type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:4530"
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'clone="HD--07-L05"
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US-09-966-880A-8 (1-198) x CF324208
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
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                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oryza sativa
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/note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: KhoI; cDNA was inserted into lambda Uni-ZAP XR yector at 5' end with EcoRI and 3' end with XhoI site. mRNA was derived from rice Histone Deacetylase overexpression line."
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/lab host="E.coli SOLR"
/clone lib="OstDAC1-overexpressing transgenic rice lambda
phage cDNA library II (HDN)"
/note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2:
XhoI; cDNA was inserted into lambda Uni_ZAP XR vector at
5' end with EcoRI and 3' end with XhoI site. mRNA was
derived from rice Histone Deacetylase overexpression
                                                                                                                                              /organism="Oryza sativa"
/mol type="mxNA"
/mol type="mxNA"
/cultivar="Nackdong"
/cultivar="Nackdong"
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/tissue type="callus"
/dev stage="proliferated callus on 2N6 media for 2 weeks"
/dab host="E.coli SOLR"
/clone lib="0sHDAC1-overexpressing transgenic rice lambda
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/tissue_type="callus"
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 RESULT 155
CF543306/c
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DB:
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Best Local Similarity:
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Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genomics and Genetics Institute, GreenGene Biotech Inc.; of Bioscience and Bioinformatics, MyongJi University
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Kim,J.S., Jun,Kim,Cheong,P.J., Kim,M.J., Le

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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Oryza sativa
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                                              CTTCATGAG 7
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Location/Qualifiers
                                                                                                                                                                                                                                                                      /note="Vector: pBluescript SK(+); Site_1: SstI; Site_2: XhoI; cDNA was inserted into lamda Uni-ZAP XR vector at 5' end with SstI and 3' end with XhoI site. Callus was induced on 2N6 media for 30 days and cultured for 36hrs on regenerated media"
                                                                                                                                                                                                                                                                                                                                                     /clone lib="Regenerated callus lambda phage cDNA library
(RCL1)"
                                                                                                                                                                                                                                                                                                                                                                              /dev_stage="proliferated callus on 2N6 media for 30 days"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Oryza sativa"
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                                                                                                                                                                                                                                                                                                                                                                                                             tissue_type="callus"
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                                           RESULT 156
CF543404/c
LOCUS
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CF543404 15 bp mRNA linear BST 22-SEP-2003 SO1468-024-029-D20-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone 024-029-D20 5-PRIME, mRNA sequence.
CF543404
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: weisshaa@mpiz-koeln.mpg.
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Carl-von-Linne Weg 10, 50829 Koeln, Germany
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Weisshaar B
ADIS DNA core facility at MPIZ
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Plate: 29 row: F column: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clome_lib="MPIZ-ADIS-024-leaf"
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Notl;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Notl;
cDNA library from sugar beet, library provided by KWS
CDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Binbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-Notl, primer sites a:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RZPD/GABI-Primary database:http://gabi.rzpd.de"
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/db_xref="taxon:161934"
/clone="024-029-F24"
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/cultivar="KWS2320 (double haploid, monogerm breeding
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/lab_host="EMDH10B"
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Menze,A., O'Brien,J., Lehrach,H.
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                                                                                                                                                      RESULT 157
R41075
                                                                                                                                                                                                                                                                                    US-09-966-880A-8 (1-198) x CF543404 (1-15)
                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                         Score:
                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                           Alignment Scores:
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                                                                                                                DEFINITION
                                       EYWORDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JOURNAL
MEDLINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DEMED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          source
                                                                                                                                                                                                                                                 193
                                                                                                                                                                                                             15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta
Spermatophyta; Magnoliophyta; eudicotyledons; core
Caryophyllales; Amaranthaceae; Beta.
1 (bases 1 to 15)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beta vulgaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CF543404.1
EST.
                                                         R41075.1 GI:798691
                                                                                            sequence.
                                                                                                            Hk082-f Adult heart,
                                                                                                                                    R41075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fingerprinting allows access to 25 plant J. 32 (5), 845-857 (2002) 22362189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and Radelof, U.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beta vulgaris
Homo sapiens (human)
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Insert Length: 15 Std Error Plate: 29 row: D column: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: weisshaa@mpiz-koeln.mpg.de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Weisshaar B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Construction of a 'unigene' cDNA clone set by oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fax: 0049221506285
                                                                                                                                                                                                             TTTCGGACG
                                                                                                                                                                                                                                                   PheArgThr 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SPG-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NcHI-T7; Note Sequencing granted in the context of the GABI-Beet project, local pri Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="MPIZ-ADIS-024-leaf"
/note="Vector: pCMVSPORT6; Site_1: Sall; Site_2: NotI;
CDNA library from sugar beet, library provided by KWS
CDNA library from sugar beet, of Sinbeck, Germany, contact:
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RZPD/GABI-Primary database:http://gabi.rzpd.de"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 orientation:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Beta vulgaris"
/mol_type="mRNA"
/cultivar="KWS2320 (double haploid, monogerm breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'db_xref="taxon:161934"
'clone="024-029-D20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lab_host="EMDH108"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       db_xref="GABI:936359"
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                                                                                              15 bp
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Matches:
Conservative:
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                                                                                                                sapiens cDNA clone k082-f,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a; Tracheophyta;
eudicots;
                                                                                                                                  EST 16-MAY-1995
                                                                                                                mRNA
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RESULT 158
HSM001764
ID HSM001
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AUTHORS
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Best Local Similarity:
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Clone from S. Wiemann, sequenced by sequencing consortium of the German No r1 sequence available This clone is available at the RZPD
                                                                                                                                                                                                                                                                                                                                                                         AL037434.1
                                                                                                                                                                                                                                                                                                                                                                                                        AL037434;
                                                                                                                                                                                                                                                                                                                                                                                                                                        HSM001764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               source
                                                                              Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                            Bloecker H.,
                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                         Homo sapiens (human)
                                                                                                                                                                                                                                                           EST; expressed sequence tag.
                                                                                                                                                                                                                                                                                          Homo sapiens mRNA;
                                                                                                                                                                                                                                                                                                                                       12-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  No.:
                                                                                                                                                                                                                                                                                                                          12-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     170 ValArgLeu 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gene expression of adult human heart as revealed by random sequencing of cDNA library
Miami Winter BioTechnol. Symp. Proc. 6, 90 (1995)
Contact: Waye Mary M.Y.
Department of Biochemistry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 15)

Waye,M.M.Y., Cheung,H.K.Y., Lam,W.Y., Law,P.T.W., Lo,A.S.Y., Lui,V.W.Y., Luk,S.C.W., Tsui,S.K.W., Tung,C.K.C., Yam,N.Y.H. Liew,C.C. and Lee,C.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The Chinese University of Hong Kong
Rm 302C, Basic Medical Science Building,
Hong Kong, Shatin, N.T., Hong Kong.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fax: 8526035123
Email: b133723@vax.csc.cuhk.hk
Seg primer: GGTGGCGACGACTCCTGGAGCC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hong Kong, Shat
Tel: 8526096874
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                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; mRNA; EST; 16
                                                                                                                                                                                                                                                                                                                        (Rel.
                                                                                                                            Boecher M., Brandt P., Mewes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="mRNA"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="k082-f"
/clone="to.coli y1090"
/clone_lib="Adult heart, Clontech"
/note="Vector: Lambda gt11; Site_1: EcoRI; Site_2: EcoRI"
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                                                sequenced by GBF within the
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                                                                                                                              Gassenhuber J.,
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at the RZPD in

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RESULT 159
HSM004270/G
ID HSM004
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DB:
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Berlin-
                                                                                                                                                                                                                                                                                                   Sequencing consortium of the German No si sequence available
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                                                                                                                                                                                                                                    This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059 Berlin-Charlottenburg, GERWANY; Email: clone@rzpd.de
                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 16 BP;
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cenburg, GERMANY; Email: clone@rzpd.de
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/organism="Homo sapiens"
/clone="DKEZp56401471"
/clone lib="564 (synonym: hfbr2). Vector pAMP1; host
X1-2blue; sites NotI + SalI"
/dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
/mol_type="mRNA"
/organism="Homo sapiens"
/organism="Homo sapiens"
/clone="DkFZp434B1612"
/clone_1lb="434 (synonym: htes3). Vector pSport1; host
DH10B; sites NotI + SalI"
                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                    db_xref="taxon:9606"
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Last updated, Version 1)
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Conservative:
Mismatches:
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AUTHORS
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AA904711/c
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VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov

Email: cgapbs-r@mail.nih.gov

This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 16)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Carger Institute, Cancer Genome Anatomy Project (CGAP);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Trace considered overall poor quality Insert Length: 1732 Std Error: 0.00 Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 1.
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                                                                                                                                                                                               /Clone_lib="Soares_NFL_T_GBC_S1"
//Clone_lib="Soares_NFL_T_GBC_S1"
//note="Organ: pooled; Vector: pT773D-Pac (Pharmacia) with
/note="Torgan: pooled; Vector: pT773D-Pac (Pharmacia) with
a modified polylinker; Site_1: Not 1; Site_2: Eco RI;
Equal amounts of plasmid DNA from three normalized
libraries (fetal lung NbHH19W, testis NHT, and B-cell
NCI_CGAP_GCB1) were mixed, and ss circles were made in
vitro. Following HAP purification, this DNA was used as
tracer in a subtractive hybridization reaction. The driver
was PCR-amplified cDNAs from pools of 5,000 clones made
from the same 3 libraries. The pools consisted of
I.M.A.G.E. clones 297480-302087, 68262-687239,
726408-728711, and 729096-731399. Subtraction by Bento
Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:1504051"
/lab_host="DH10B"
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/tissue_type="testis"
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AA939272/c
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oq31b06.s1 NCI CGAP GC4 Home sapiens CDNA clone IMAGE:1587923
similar to 5W:CR34 HUMAN Q01955 POCOCLAGEN ALPHA 3(IV) CHAIN
PRECURSOR. ;contains OFR.b3 MSR1 repetitive element ;, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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1 (bases 1 to 16)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1997)
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                                      GlyAsnPro 102
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/note="Vector: pT713D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from 3 pooled polylinker; 1st strand then primed with a Not I germ cell tumors, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tissue_type="pooled germ cell tumors"
(lab_host="DH108"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone="IMAGE:1587923"
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AA953804/c
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                                                                                                                                    RESULT 163
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AA968729 16 bp mRNA linear EST 27-AUG-
Or69h11.s1 NCI_CGAP_GC3 Homo sapiens cDNA clone IMAGE:1601157 2
similar to SW:PRPE HUMAN P02811 BASIC PROLINE-RICH PEPTIDE P-E
;contains element MSR1 repetitive element ; mRNA sequence.
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Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Trace considered overall poor quality Insert Length: 434 Std Error: 0.00 Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 1.
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CONA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Cente
Clone distribution: NOI-CGAP clone distribution information can
found through the I.M.A.G.B. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
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National Cancer Institute, Cancer Genome Anatomy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone lib="NCI_CGAP_Lu5"
/note="Organ: lung; Vector: pT7T3D-Pac (Pharmacia) with a /note="Organ: lung; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; lst strand cDNA was prepared from neuroendoorine lung carcinoid, and was then primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and Cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tissue_type="carcinoid"
lab_host="DH10B"
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db_xref="taxon:9606"
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                                                                                                                                                                                                                         AA968729/c
                                                                                                                                                                                                                                               RESULT 164
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or69hl1.51 NCI_CGAP_GC3 Homo sapiens cDNA clone IMAGE:1601157 :
similar to SW:FRPE HUMAN PO2811 BASIC PROLINE-RICH PEPTIDE P-E
contains element MSR1 repetitive element ; mRNA sequence.
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Unpublished (1997)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 16)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mational Cancer Institute, Cancer Genome Anatomy Project (CGAP)
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                                                           Homo sapiens (human)
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/note="Vector: pT773D-Pac (Pharmacia) with a modified polylinker; lst strand cDNA was prepared from 3 pooled polylinker; lst strand cDNA was prepared with a Not I oligo(dr) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and Cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is not normalized. Library was
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/tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
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/mol_type="mRNA"
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Contact: Robert Strausberg, Ph.D
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Query Match:
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Insert Length; 514 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 1.
                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 16)

NCI/NINDS-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project
                                                                                                                                                                                                                                                                                                                                                              element ;, mRNA sequence.
AI094839
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CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
(CGAP/BTGAP), Tumo
Unpublished (1998)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/note="Vector: p17/13D-Pac (Pharmacia) with a modified
/note="Vector: p17/13D-Pac (Pharmacia) with a modified
polylinker; lst strand cDNA was prepared from 3 pooled
perm cell tumors, and was then primed with a Not I
oligo(dT) primer. Double-stranded cDNA was ligated to Eco
RI adaptors (Pharmacia), digested with Not I and Cloned
into the Not I and Eco RI sites of the modified p17/13
vector. Library is not normalized. Library was
vector. Library is not normalized. Library was
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/lab_host="DH10B"
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Ph.D.
               Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 543 Std Error: 0.00
                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 16)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                16 bp mRNA linear EST 28-OCT-1998 qd36b06.x1 Soares fetal heart NbH19W Homo sapiens cDNA clone IMAGE:1731539 3' similar to SW:SN22 HUMAN P51531 POSSIBLE GLOBAL TRANSCRIPTION ACTIVATOR SNF2L2 ;, mRNA sequence.
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DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.B. Consortium/LLNL at:
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                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                    Unpublished (1997)
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Procurement: David N. Louis, M.D., Myrna R. Rosenfeld M.D.,
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/lab_host="DH10B"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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FEATURES

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AI208066
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                                                                                                                                                                                                                                                 JOURNAL
                                                                                                                                                                                                                                                                               TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                       16 bp mRNA linear EST 29-NOV-1998 qg46a09.xl Soares testis NHT Homo sapiens cDNA clone IMAGS:1838200 3' similar to WP:M7.1 CE03482 LET-70: UBIQUITIN-CONJUGATING ENZYME E2-17 KD ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38 SerAlaThr 40
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                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
cDNA_Library_Preparation: M. Bento Soares, Ph.D.,
                                               Trace considered overall poor quality Insert Length: 618 Std Error: 0.00
                                                                                                                                                                                                                                                                            NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 16)
                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                              Insert Length: 618 Std Err
Seq primer: -40UP from Gibco
                                                                                                            CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E, Consortium/LLNL at:
                                                                                                                                                                                                                                               Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                             AI208066.1
                                                                                               www-bio.llnl.gov/bbrp/image/image.html
quality sequence stop: 1.
Location/Qualifiers
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:1731539"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ocation/Qualifiers
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SOURCE
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Best Local Similarity:
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Smith College Department of Biological Sciences
Department of Biological Sciences, Clark Science Center, Smith
College, Northampton, MA, 01063, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Williams, S.A., Lizotte-Waniewski, M., Laney, S. and Lustigman, S. Genes expressed in molting L3 larvae of Onchocerca volvulus Unpublished (1997)
Contact: Steven A. Williams
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AI603831

16 bp mRNA linear EST 21-APR-199
SWOV3WCAM27A07SK Onchocerca volvulus molting L3 larva cDNA
(S196MLW-OvmL3) Onchocerca volvulus cDNA clone SWOV3WCAM27A07 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Onchocerca volvulus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AI603831
AI603831.1 GI:4612980
                                                                                                                                                                                                                                                                                        Seq primer: pBluescript SK.
                                                                                                                                                                                                                                                                                                                                                        Tel: 4135853826
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA sequence.
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                                                                                                                                                                                                                                                                                                             Email: genome@smith.edu
                                                                                                                                                                                                                                                                                                                                      Fax: 4135853786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Onchocercidae; Onchocerca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Nematoda; Chromadorea; Spirurida; Filarioidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ArgValThr 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (bases 1 to 16)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone libe"Soares testis NHT"
/clone libe"Soares testis NHT"
/note="Vector: pT7T3D-Pac (Pharmacia) with a modified
polylinker; Site 1: Not I; Site 2: Eco RI; 1st strand cDNA
was prepared from mRNA obtained from Clontech
Laboratories, Inc., and primed with a Not I - oligo(dT)
                                                                                                                                                     /mol_type="mRNA"
/strain="Kumba, Cameroons"
/db_xref="taxon:6282"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
note="Vector: Lambda Uni-ZAP XR; Site_1: Eco RI; Site_2:
Tho I; Filarial nematode parasite of humans. Third-stage
                                                                                   dev_stage="molting L3"

lab_host="XL1-Blue MRF'"
                                                                                                                                                                                                                                                                  location/Qualifiers
                                             SL96MLW-OVML3)"
                                                               clone_lib="Onchocerca volvulus molting L3 larva cDNA
                                                                                                                                   clone="SWOv3MCAM27A07"
                                                                                                                                                                                                                    organism="Onchocerca volvulus"
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Score:

Pred. No.:

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RESULT 168

AI603831

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DEFINITION

REFERENCE AUTHORS TITLE

VERSION KEYWORDS

ACCESSION

COMMENT

FEATURES

Location/Qualifiers

JOURNAL

larvae, L3, were isolated from infected black flies in Camerson (forest strain). The L3 were cultured in 20% PCS in IMDM+ NCTC 135 and collected after day 1, 2, or 3 in culture. L3 of 0. volvalus molt to fourth-stage larvae by day 5 in culture. mRNA was isolated from approximately 6000 molting larvae (mL3),2000 larvae from day 1, 2 or 3 in culture, and converted to double-stranded cDNA using reverse transcriptase and oligo(dT) followed by RNase H and DNA pol 1. The library was constructed in the lambda Uni-Zap XR vector and has 1 x 10E6 independent recombinants and the average insert size is -1200 bp. The library was constructed by Sara Lustigman and Michelle Lizotte-Waniewski in the Laboratory of Dr. S. A. Williams. The library is available from Dr. Sara Lustigman (email: slustigm@nybc.org)."

ORIGIN

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REFERENCE
AUTHORS
TITLE
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ORGANISM
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AW248540
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                                                                                   Contact: Robert Straubberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E.
Consortium (LLNI) DNA Sequencing by: Berkeley MGC sequencing
project Clone distribution: MGC clone distribution information can
be found through the I.M.A.G.E. Consortium/LLNI at:

www-bio.llni.gov/bbrp/image/image.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center: Vector
Trimming: cross match from University of Washingtion Genome Center
PHRAP suite. Poly-T Identification: patMatch.pl from Berkeley
Drosophila Genome Project. University of Washingtion Genome Center:
http://www.genome.washington.edu Low Quality Sequence: 15
contiguous PHRED high quality bases following vector sequence. Very
Low Quality Sequence: Trace file contained 16 contiguous distinct
peaks following vector sequence. Polyadenylation: Based upon the
presence of a XhoI site followed by a run of 14 or more T residues
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                         polyadenylated.
Plate: "' ~ .
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 16)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1999)
Other_ESTs: 2820844.5prime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AW248540 16 bp menA linear ESI V/-VAN-2VV 2820844.3prime NIH_MGC_7 Homo sapiens cDNA clone IMAGE:2820844 3',
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                      .ate: LLCM5 row: E column:
quality sequence
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Matches:
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                                                                                   source
                                                                                                                                                                                                      Other ESTs: 282266.5prime
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue procurement: DCTD/DTP cDNA Library Preparation: Ling
Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E.
Consortium (LLNL) DNA Sequencing by: Berkeley MGC sequencing
project Clone distribution: MGC clone distribution information can
be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center.
Trimming: cross match from University of Washingtion Genome Center
PHRED high quality bases following vector Sequence: 9 contiguous
PHRED high quality bases following vector sequence. Very Low
Quality Sequence: Trace file contained 16 contiguous distinct peaks
following vector sequence. Polyadenylation: Based upon the presence
beginning of the sequence of the cortained at the presence of a Xhol site followed by a run of 14 or more T residues at the
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1999)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                               beginning of the sequence, Plate: LLCM8 row: P colu
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                                                       quality sequence stop: 9.

Location/Qualifiers

Listinct

List DNA in or more T residues at the pre-

List Town or P column: 12

Quality sequence stop: 9.

Location/Qualifiers

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/tissue_type="small cell carcinoma"
/tissue_type="small cell carcinoma"
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cloned into EccRIXhoI sites using the following 5'
edaptor: GGCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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VERSION **EYWORDS**

FEATURES

Alignment Scores: Pred. No.:

3.39e+06 3.00

Length: Matches:

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Score:

ORIGIN

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REFERENCE
AUTHORS
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BG897738
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Osteoarthr. Cartil. 9 (7), 641-653 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kumar,S., Connor,J.R., Dodds,R.A., Halsey,W., Van Horn,M., Mac
Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen,M.
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1 (bases 1 to 16)
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BG897738
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                                                                                                                                                                                                                                                         Fax: 610-270-5598
                                                                                                                                                                                                                                                                                 Tel: 610-270-7245
                                                                                                                                                                                                                                                                                                                           GlaxoSmithKline
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/cell line="MGC3"
/lab_host="DH10B (phage-resistant)"
/clome_lib="NIH_MGC 7"
/clome_lib="NIH_MGC 7"
/clome_lib="NIH_MGC 7"
/clome_lib="NIH_MGC 7"
/clome_lib="NIH_MGC 7"
/clome_lib="NIH_MGC 7"

EcoRI; CDNA made by oligo-dT priming. Directionally clomed into EcoRI/KhoI sites using the following 5'
adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
                                                                                                                                                                                                                                     sanjay
/tissue_type="cartilage"
/lab host="E.coli DHIO B"
/clone_lib="HOA (Human Osteoarthritic Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: No
Directional"
                                                                                                       mol_type="mRNA"
db_xref="taxon:9606"
                                                                                                                                               organism="Homo sapiens"
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RESULT 173
BG926060/c
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    HNC23-1-E1.R . sequence. BG926060
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 16)
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BG900981.1 GI:14311230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: sanjay_kumar-1@gsk.com
Seq primer: T7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fax: 610-270-5598
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 709 Swedeland Road,
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/lab host="E.coli DHI0 B"
/clone_lib="HOA (Human Osteoarthritic Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: Not
Directional"
                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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Cartilage)
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                                linear EST 06-NOV-2001 Homo sapiens cDNA, mRNA
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BM397104/c
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                                                                                                                                                                                                                                                                                                                                                                                                                  34 Lysargarg
                                                                                                                                                                          Tetrahymena thermophila
Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota, Alveolatta, Ciliophora, Oligohymenophorea,
Hymenostomatida, Tetrahymenina, Tetrahymena.
                                                                                                                                                                                                                                                                                       16 bp mRNA linear EST 17 5009-0-28-H09.t.2 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
Contact: Turkewitz AP
Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 606
Tel: 773 702 4374
Fax: 773 702 3172
                                                                                               EST from Tetrahymena thermophila, 
Unpublished (2002)
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                                                                                                                            1 (bases 1 to 16)
Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E.,
Frankel,J. and Klobutcher,L.
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Tel: 610-270-7245
Fax: 610-270-5598
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Osteoarthr. Cartil. 9 (7), 641-653 (2001)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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EST.
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Seq primer: T7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue type="cartilage"
/lab_host="E.coli DH10 B"
/lab_host="E.coli DH10 B"
/clone lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: Sall; Site_2: NotI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mol_type="mRNA"
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                                Chicago, IL 60637, USA
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                                                                                                              growing cells
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RESULT 175
BM399406
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Turkewitz,A.P., Karrer,K.M.,
Frankel,J. and Klobutcher,L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16 bp mRNA linear EST 17 5009-0-57-D08.t.2 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                          Tel: 773 702 4374 Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                       Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 60637, USA
                                                                                                                                                                                                                                                                                                                          Email: apturkew@midway.uchicago
                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Turkewitz AP
                                                                                                                                                                                                                                                                                                                                                                                                                                      EST from Tetrahymena thermophila, strain CU428.1, growing cells Unpublished (2002)
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/clone lib="chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (20
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                                                                          /clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/strain="CU428.1"
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                              Matches:
Conservative:
Mismatches:
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BM817126/c
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BM401358/c
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VERSION
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                       AUTHORS
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           Hordeum vulgare subsp. vulgare
Hordeum vulgare subsp. vulgare
Eukaryota; Viridiplantae; Streptophyta; Em
Spermatophyta; Magnoliophyta; Liliopsida;
Pooideae; Triticeae; Hordeum.

1 (bases 1 to 16)
0zturk,N.Z., Michalowski,C.B., Brazille,S.
                                                                                                                                                                                          HC02D04_T3.ab1 HC Hordeum vulgare HC02D04_T3.ab1, mRNA sequence.
BM817126
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University of Chicago
920 E. 58th Street, Chicago, IL 6061
Tel: 773 702 4374
Fax: 773 702 3172
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Turkewitz, A.P., Karrer, K.M., Jahn, C., Orias, E., Kirk, K.E., Frankel, J. and Klobutcher, L.
BST from Tetrahymena thermophila, strain CU428.1, growing cells Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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Ozturk, N.Z., Michalowski, C.B., Brazille, S., Borchert, C., Palacio, C., Normand, C., Murphy, C., Kelley, R., Sant, S.A.,
                                                                                                                                                                        BM817126.1 GI:19153140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: apturkew@midway.uchicago.
Seq primer: T3.
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/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/strain="CU428.1"
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S011421-024-001-L05-SP6R MPIZ-ADIS-024-inflorescence
CDNA Clone 024-001-L05 5-PRIME, mRNA sequence.
                                                                                                                                     Contact: Weisshaar B
ADIS DNA core facility at MPIZ
AMAx-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
                             Email: weisshaa@mpiz-koeln.mpg.de
Insert Length: 16 Std Error: 0.00
Plate: 1 row: L column: 05
Seq primer: SP6r; ATTTAGGTGACACTATAGAAGA.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                             Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H. and Radelof,U.
                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Caryophyllales; Maranthaceae; Beta.

1 (bases 1 to 16)
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                                                                                                                                                                                                                                                   Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet ge Plant J. 32 (5), 845-857 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beta vulgaris
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beta vulgaris
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/organism="Beta vulgaris|
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone lib="HC"
/note="6 and 10 hour drought stress by placing
/notst paper (75% rel. humidity) in light"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /db_xref="taxon:112509"
/clone="HC02D04_T3.ab1"
/tissue_type="Root"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dev_stage="3 week old"
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strain="cv tokak"
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Query Match:
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
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E012394-024-013-N21-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone
                                                                                                                                                                                                                                                                                                                                       Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                         Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,
Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Leh
                                                                                                                                                                                                                                                                                                                                                                                                                                                              BQ586020
BQ586020.1 GI:26115602
                                                                                                     Insert Length: 16 Std Error: 0.00
Plate: 13 row: N column: 21
Seq primer: SP6; CATACGATTTAGGTGACACTATAG
                                                                                                                                                                    Email: weisshaa@mpiz-koeln.mpg.de
                                                                                                                                                                                            Fax: 00492215062851
                                                                                                                                                                                                                                                                          Contact: Weisshaar B
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/lab_host="EMDHIOB"
/clone_lib="MPIZ-ADIS-024-inflorescence"
/clone_vector: pCMVSPORT6; Site_1: Sall; Site_2: NotI;
/note="Vector: pCMVSPORT6; Site_1: Sall; Site_2: NotI;
/cDNA_library_from sugar_beet, library_provided_by_KWS
Kleinwanzlebener_Saatzucht_AG_Einbeck, Germany, contact:
b.schulz@kws.de; cloning_sites_Sall-NotI, primer_sites_ai
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                       /organism="Beta vulgaris"
/mol_type="mRNA"
                                                                                      location/Qualifiers
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/db_xref="taxon:161934"
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/cultivar="KWS2320 (double haploid, monogerm breeding
cultivar="KWS2320 (double haploid, monogerm breeding
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xref="GABI:186842"

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1 (bases 1 to 16)

Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,

Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.

and Radelof, U.

Construction of a 'unigene' cDNA clone set by oligonucleotide

fingerprinting allows access to 25 000 potential sugar beet genes

Plant J. 32 (5), 845-857 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Max-Planck-Institute for Plant Breeding Research Carl-von-Linne Weg 10, 50829 Koeln, Germany Fax: 00492215062851
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ADIS DNA core facility at MPIZ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: weisshaa@mpiz-koeln.mpg.de
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ert Length: 16 Std Error: 0.00
te: 13 row: C column: 19
primer: SP6; CATACGATTAGGTGACACTATAG
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/clone lib="MMPIZ-ADIS-024-leaf"
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Notl;
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/note="Vector: pcmvSpORT6; Site 1: Sall; Site 2: Notl;
/note="Vector: pcmvSpORT6; Site 3: Notl; Site
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/clone="024-013-N21"
/clone="024-013-C19"
/tissue_type="leaf"
/lab_host="EMDH108"
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1 (bases 1 to 16)
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
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Plate: 10 row: M column: 01
Seq primer: SP6; CATACGATTTAGGTGACACTATAG
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cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact
b.schulz@kws.de; cloning sites Sall-Notl, primer sites
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/clome_lib="MPIZ-ADIS-024-leaf"
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cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contac
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E012340w-024-010-M01-SP6 MPIZ-ADIS-024-leaf
clone 024-010-M01 5-PRIME, mRNA sequence.
BQ587767
BQ587767.1 GI:26117349
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Max-Planck-Institute for Plant Breeding Research
Carl-von_Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
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1 (bases 1 to 16)
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                                                                    /note=TVector: pCMVSPORT6; Site 1: SalI; Site 2: NotI; CDNA library from sugar beet, library provided by KwS Kleinwanzlebener Sastzucht AG Einbeck, Germany, contact: b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
SP6-SalI-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7;
Sequencing granted in the context of the GABI-Beet
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
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Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
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/DNA library from sugar beet, library provided by KMS
cDNA library from sugar beet, library provided by KMS
Kleinwanzlebener Saatzucht AG Binbeck, Germany, contact
b.schulz@kws.de; cloning sites Sall-NotI, primer sites
Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database:http://gabi.rzpd.de"
                                                                   SPG-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Sequencing granted in the context of the GABI-Beet
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                          lab_host="EMDH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Beta vulgaris"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
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Plate: 9 row: A column: 19
Seg primer: SP6; CATACGATTTAGGTGACACTATAG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: weisshaa@mpiz-koeln.mpg.de
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                                                                                                                                                                                                                   /clone lib="MPIZ-ADIS-024-leaf"
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not!;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not!;
cDNA library from sugar beet, library provided by KWS
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-Not!, primer sites and
                                                                                              SP6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database:http://gabi.rzpd.de"
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                                                                                                                                                                                                                                                                                                                        tissue_type="leaf"
/lab_host="EMDH10B"
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/clone="024-009-A19"
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
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Sp6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de"
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/cDNA library from sugar beet, library provided by KWS
CDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact
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/mol_type="mRNA"
/cultivar="KW82320 (double haploid, monogerm breeding
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                                                                                                                              Similarity:
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16 bp mRNA linear EST 06-DEC-20
E012562-024-015-N03-SP6 MPIZ-ADIS-024-storage root Beta vulgaris
cDNA clone 024-015-N03 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Amaranthaceae; Beta.
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EST.
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Insert Length: 16 Std Error: 0.00
Plate: 15 row: N column: 03
Seq primer: SP6; CATACGATTTAGGTGACACTATAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="storage root"
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/clone_lib="MPIZ-ADIS-024-storage root"
/clone_tib="MPIZ-ADIS-024-storage root"
/note="Vector: pCMVSPORT6; Site_1: SalI; Site_2: NotI;
/cDNA_library_from sugar beet, library_provided_by_KWS
Kleinwanzlebener_Sastzucht_AG_Einbeck, Germany, contact:
b.schulz@kws.de; cloning_sites_SalI-NotI, primer_sites_and
                                                                                                                                                                                                                                                                                                 Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by
                                                                                                                                                                                                                                                                                                                                                               SP6-SalI-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Note:
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/mol_type="mRNA"
/cultivar="KWS2320 (double
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/clone="024-015-N03"
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Best Local Similarity:
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                                                          HDA1--03-G11.g1 OsHDAC1-overexpressing transgenic rice lambda cDNA library I (HDA1) Oryza sativa cDNA clone HDA1--03-G11, mF
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Genomics and Genetics Institute, GreenGene Biotech Inc.;
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ABF1--03-B14.g1 ABF3-overexpressing transgenic rice lambda phage cDNA library (ABF1) Oryza sativa cDNA clone ABF1--03-B14, mRNA
CF306313.1
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Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1. (bases 1 to 16)
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                                                                                                  CF306313
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    Location/Qualifiers
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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/db_xref="taxon:4530"
/clone="ABF1--03-B14"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="ABF3-overexpressing transgenic rice lambda phage cDNA library (ABF1)"
note="Vector: pBluescript SK(+); Site_1: EcoRI; Site_2: XhoI; Leaf was dried for 2hrs. cDNA was inserted into lamda Uni-ZAP XR vector at 5' end with EcoRI and 3' end with XhoI site. mRNA was prepared from ABA-responsive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              element binding transcription factor 3 overexpression
line."
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/lab_host="E.coli SOLR"
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                                                                                                linear
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Oryza sativa

REFERENCE AUTHORS

COMMENT

TITLE

FEATURES

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ORGANISM
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CF307345/c
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Genomics and Genetics Institute, Greendene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
Yongin, Kyeenggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Genomics and Genetics Institute, (
of Bioscience and Bioinformatics,
Yongin, Kyeenggi, Korea
Tel: 82 31 330 6193
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Oryza sativa
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Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                                     Contact: Nahm B.H.
                                                                                                                                                                                                                                                                (bases 1 to 16)
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: XhoI; Callus was treated with ABA(20um) for Ihour. cDNA was inserted into lambds Uni-ZAP XR vector at 5' end with EcoRI and 3' end with XhoI site. mRNA was derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rice Histone Deacetylase overexpression line."
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phage cDNA library I (HDA1)"
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/cultivar="Nackdong"
/db_xref="taxon:4530"
/clone="HDA1--03-G11"
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Matches:
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, MyongJi University
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                                                                                    Inc.; Division
   THE HEAR COCOCCEPTED AND COCCEPTED AND COCCE
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Eutheria; I
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                                                                                                                                                                       Clone from S. Wiemann, sequenced by LMU within the cDNA sequencing consortium of the German Genome Project No sl sequence available This clone is available at the RZPD in Berlin
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Location/Qualifiers
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se contact the RZPD: Ressourcenzentrum,
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                                                  /mol_type="mRNA"
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DH10B; sites NotI + SalI"
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/clone lib="434 (symonym: htes3). Vector pSport1; host DH10B; sites Not! + Sall"
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                                                                                                                                                                                                         /organism="Homo sapiens"
/clone="DKFZp434K1222"
/clone=lib="434 (synonym: htes3). Vector pSport1;
DH10B; sites Not1 + SalI"
/dev_srage='adult"
/tissue_type="testis"
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/mol_type="mRNA"
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleosto Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 17)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
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1 (bases 1 to 17)

Brenner,S., Williams,S.R., Vermass,E.H., Storck,T., Moon,K., McCollum,C., Mao,J.I., Kirchner,J.J., Eletr,S., DuBridge,R.B., Burcham,T. and Albrecht,G.

Eurcham,T. and Albrecht,G.

In vitro cloning of complex mixtures of DNA on microbeads: Physical separation of differentially expressed cDNAs

Proc. Natl. Acad. Sci. U.S.A. 97 (4), 1665-1670 (2000)
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                                                                                                                   Homo sapiens
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25861 Industrial Blvd., Hayward, CA 94545, USA
                                                                                                                                     Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: timb@lynxgen.com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Burcham TS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20144098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               quality sequence stop: 17.
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510 670 9302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone lib="DNC15"
/note="Vector: pCR2.1; Cloning of PCR products from micro-beads carrying 3' end of down-regulated cDNA.cells non-induced (treated with DMSO only)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo mapiens"

/mol_type="mRNA"

/db xref="taxon:9606"

/cell_type="monocyctic leukemia"

/cell_line="THP-1 (TIB-202)"

/clone lib="DNC15"
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100.00%
1.52%
                                                                                                                                                                            GI:6589521
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                                                                                                                                                                                                                               17 bp mRNA linear EST 07-JAN-
Homo sapiens cDNA clone IMAGE:2821879
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Conservative:
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AGE:2821879 3',
                                                                              Euteleostomi;
Homo.
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SOURCE
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AW246893/c
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DB:
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Best Local Similarity:
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Consortium (LLML) DNA Sequencing by: Berkeley MGC sequencing
project Clone distribution: MGC clone distribution information can
be found through the I.M.A.G.E. Consortium/LLML at:
www-bio.llnl.gov/bbrp/image/image.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center. Vector
Trimming: cross match from University of Washingtion Genome Center
PHRAP suite. Poly-T Identification: patMatch.pl from Berkeley
Drosophila Genome. Project: University of Washingtion Genome Center:
http://www.genome.washington.edu Low Quality Sequence: 13
contiguous PHRED high quality bases following vector sequence. Polyadenylation: Based upon the
presence of a XhoI site followed by a run of 14 or more T residues
and the beginning of the sequence, this cDNA insert was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15 PheLysAsn 17
Unpublished (1999)
Other_ESTs: 2822293.3prime
                                               NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health,
                                                                                                                                                                                        Homo sapiens
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Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
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Contact: Robert Strausberg,
                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 17)
                                                                                                                                                                                                                                                                                                                         mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                2822293.5prime NIH_MGC_7 Homo sapiens cDNA clone IMAGE:2822293 5',
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTTAAAAAC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /cell līne="MGC3"
/lab host="DH10B (phage-resistant)"
/lab host="DH10B (phage-resistant)"
/clone lib="NH1MGC 7"
/clone lib="NH1MGC 7"
/note="Organ: lung; Vector: pOTB7; Site 1: XhoI; Site 2: /note="Organ: lung; Vector: pOTB7; Site 1: XhoI; Site 2: /note="Organ: lung; Vector: pOTB7; Site 1: XhoI; Site 2: /note="Organ: lung; Vector: potential following 5"
cloned into EccRI/XhoI sites using the following 5"
adaptor: GCCACGAG(G). Size-selected >500bg for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
                                                                                                                                                                                                                    sapiens (human)
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/tissue_type="small cell carcinoma"
/cell_line="MGC3"
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                                                        Mammalian Gene Collection (MGC)
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RESULT 196
AW247949
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Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E.
Consortium (LLML) DNA Sequencing by: Berkeley MGC sequencing
project Clone distribution: MGC clone distribution information can
be found through the I.M.A.G.E. Consortium/LLML at:
www-bio.llnl.gov/bbrp/image/image.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center. Vector
Trimming: cross match from University of Washingtion Genome Center
PHRAP suite. Poly-T Identification: patkMatch.pl from Berkeley
Drosophila Genome Project. University of Washingtion Genome Center:
http://www.genome.washington.edu Low Quality Sequence: 7 contiguous
PHRED high quality bases following vector sequence. Very Low
Quality Sequence: Trace file contained 17 contiguous distinct peaks
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                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 17)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)

Other_ESTs: 2820605.5prime
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E.
                                                                                       Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 mRNA sequence.
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/lab host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_7"
/clone_lib="NIH_MGC_7"
/clone_lib="NIH_MGC_7"
/clone_lib="NIH_MGC_7"
/clone_lib="NiH_MGC_7"
/clone_lib="NiH_MGC_7"
/cloned into EccRI/XhoI sites using the following 5'
adaptor: GGCAGCAG(G). Size-selected >500bg for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-CDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
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Location/Qualifiers
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/tissue_type="small cell carcinoma"
/cell_line="MGC3"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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Indels:
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SOURCE
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BG926068/c
LOCUS
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                                      JOURNAL
MEDLINE
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                                                                                                                    sequenced tags (ESTs) each from adult hu osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (200)
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                                                                                                                                                                                                                                                                                                                                    Kumar, S., Connor, J.R., Dodds, R.A., Halsey, W., Van Horn, M., Mao, J.,
Sathe, G., Mui, P., Agarwal, P., Badger, A.M., Lee, J.C., Gowen, M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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Contact: Sanjay Kumar
                                      11597177
                                                                                                                                                                                                           Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and
                                                                                                                                                                                                                                                                                                 Lark, M.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BG926068.1 GI:14320591
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//lab host="DH108 (phage-resistant)"
//lab host="DH108 (phage-resistant)"
//lab host="DH108 (phage-resistant)"
//lab host="DH18 MGC 7"
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/cell_line="MGC3"
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/clone="IMAGE:2820605"
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mol_type="mRNA"
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Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E.,
Frankel,J. and Klobutcher,L.
BST from Tetrahymena thermophila, strain CU428.1, growing cells
Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                   Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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Tel: 610-270-7245
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Seq primer: T7.
                                                                                                                                                                                                                              Email: apturkew@midway.uchicago.
                                                                                                                                                                                                                                            Tel: 773 702 4374 Fax: 773 702 3172
                                                                                                                                                                                                                                                                                            Contact: Turkewitz AP
Molecular Genetics and Cell Biology
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                                                                /tissue_type="cartilage"
/lab_host="E.col1 DH10 B"
/clone_lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Sit
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/mol_type="mRNA"
/db_xref="taxon:9606"
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/mol_type="mRNA"
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Tetrahymena thermophila
Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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Percent Similarity:
Best Local Similarity:
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Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
                                   5009-0-84-D08.t.1 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
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Molecular Genetics and Cell Biology
University of Chicago
920 B. 58th Street, Chicago, IL 606
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Turkewitz, A.P., Karrer, K.M.,
Frankel, J. and Klobutcher, L.
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5009-0-19-G03.t.1 Chilcoat/Turkewitz cDNA (large fraction)
Tetrahymena thermophila cDNA, mRNA sequence.
BM401224.1 GI:18201277
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                                                                                                                                                                                     ArgValThr 79
                                                                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (20
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/mol_type="mRNA"
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Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E.,
Frankel,J. and Klobutcher,L.
EST from Tetrahymena thermophila, strain CU428.1, growing cells
                                                                                                                                        Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                     Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Amaranthaceae; Beta.

1 (Dases 1 to 17)

Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2002)
Contact: Turkewitz AP
Molecular Genetics and Cell Biology
                                                                 Email: weisshaa@mpiz-koeln.mpg.de
Insert Length: 17 Std Error: 0.00
Plate: 9 row: 0 column: 24
Seq primer: SP6; CATACGATTTAGGTGACACTATAG.
                                                                                                                                                                                                                                                                 Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
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920 E. 58th Street, Chicago,
Tel: 773 702 4374
Fax: 773 702 3172
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Seq primer: T3.
1. .17
/organism="Beta vulgaris"
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/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                Location/Qualifiers
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yaris cDNA clone
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AUTHORS
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US-09-966-880A-8 (1-198) x BQ587868 (1-17)
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                                                                                                                                                                                                                                                                                                              Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet ge Plant J. 32 (5), 845-857 (2002)
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[[bases 1 to 17]]

[bases 1 to 17]
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BQ590447
BQ590447.1 GI:26120030
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                                                                                                                                                        Email: weisshaa@mpiz-koeln.mpg.de
Insert Length: 17 Std Error: 0.00
Plate: 19 row: 0 column: 05
Seq primer: SP6; CATACGATTTAGGTGACACTATAG
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b.schulz@kws.de; cloning sites Sall-NotI, primer sites
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                      cultivar="KWS2320 (double haploid, monogerm breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="GABI:184994"
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/cultivar="KWS2320 (double haploid, monogerm breeding
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                                                                                                                                                                                                                                                                                           Carl-von-Linse Weg 10, 50829 Koeln, Germany Fax: 00492215062851
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Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,
Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,
and Radelof,U.
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te: 26 row: I column: 23
primer: SP6; CATACGATTAGGTGACACTATAG.
Location/Qualifiers
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/note="Vector: pCMVSPORT6; Site 1: Sal; Site 2: Not!;
cDNA library from sugar beet, library provided by KWS
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
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/lab_host="EMDH10B"
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'clone="024-019-005"
                                                            db_xref="taxon:161934"
                                                                                                                      cultivar="KWS2320 (double haploid, monogerm breeding
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RESULT 203 BQ593528 LOCUS

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SOURCE ORGANISM VERSION KEYWORDS

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3.61e+06 3.00 100.00% 100.00% 1.52%

Length: Matches: Conservative:

Mismatches: Indels:

0 0 0 3 17

JOURNAL MEDLINE TITLE

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/note="Vector: pCMVSPORT6; Site 1: Sal; Site 2: Not!;
cDNA library from sugar beet, library provided by KMS
CDNA library from Sugar beet, library provided by KMS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites an
                                    Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by
Prof. Christian Jung; Sequence submission manag
RZPD/GABI-Primary database: http://gabi.rzpd.de<sup>m</sup>
                                                                                                                                                  SP6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7;
Sequencing granted in the context of the GABI-Beet
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US-09-966-880A-8 (1-198) x BQ593528
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                                                                                                                                                                                                   Contact: Lambrecht M
The Arabidopsis Information Resource
Carnegie Institution of Washington, Der
260 Panama Street, Stanford, CA 94305,
Tel: 1 650 325 1521 x 251
Fax: 1 650 325 3748
                                                                                                                                                                                                                                                                                                                                                                                          Clarke,B., Lambrecht,M. and Rhee,S.Y.
Arabidopsis genomic information for interpreting wheat
                                                                                                                                                                                                                                                                                                                                                                                                                            Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magmoliophyta, Liliopsida, Poales, Poaceae, Pooldeae, Triticeae, Triticum.

1 (base 1 to 17)
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                                                                                                                                                                                    rhee@acoma.stanford.edu.
                                                                                                                                                                                                                                                                                                                                                         Integr. Genomics 3 (1-2), 33-38 (2003)
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/db_xref="taxon:4565"
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/clone_lib="wheat_EST_endosperm_library"
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Oryza sativa
Oryza sativa
Oryza sativa
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                  17 bp mRNA linear EST 15-AUG-2003
7LEAF--03-N22.gl Rice leaf plasmid cDNA library II (7LEAF) Oryza
sativa cDNA clone 7LEAF--03-N22, mRNA sequence.
CF299737
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Fax: 1 650 325 3748
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BRY_1399 wheat EST endosperm
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="endosperm"
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)

Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                1 (bases 1 to 17)

Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

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Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com,
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Tel: 82 31 330 61
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/cultivar="Nackdong"
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/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
                                                                                               location/Qualifiers
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/clone="7LEAF--03-N22"
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/mol_type="mRNA"
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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cDNA library
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Location/Qualifiers
                                                                                                                                                                                                                                                                                    /mol_type="mRNA"
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phage cDNA library I (HDA1)"
/note="Vector: pBluescript SK(+); Site_1: EcoRI; Site_2:
XhoI; Callus was treated with ABA(20um) for lhour. cDNA
was inserted into lambda Uni-ZAP XR vector at 5' end with
EcoRI and 3' end with XhoI site. mRNA was derived from
rice Histone Deacetylase overexpression line."
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/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with_oligoribonuclectides and then used as templates for
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CF921142 17 bp mRNA gmrhRww3-05_H07_1_049 Soybean root hair s gmrhRww3 Glycine max cDNA, mRNA sequence.
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Fax: 82 31 321 6355
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REFERENCE AUTHORS

COMMENT

TITLE JOURNAL

FEATURES

Score: Alignment ORIGIN

Pred. No.:

VERSION KEYWORDS

ACCESSION

SOURCE

ORGANISM

RESULT 208 CF306383

DEFINITION

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ORIGIN

Alignment

No.:

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1 (bases 1 to 17)
1 (bases 1 to 17)
2 (bases 1, Kim, K.M., Cheong, P.J., Kim, M.J., Le Kim, J.S., Jun, K.M., Kim, Y.-K. and Nahm, B.H. Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Large-scale Sequencing Analysis 
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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17 bp mRNA linear EST 18-AUG-2003 RCL1--04-J13.gl Regenerated callus lambda phage cDNA library (RCL1) Oryza sativa cDNA clone RCL1--04-J13, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: pBluescript SK(+); Site_1: StI; Site_2: XhoI; cDNA was inserted into lamda Uni-ZAP XR vector at 5' end with SsII and 3' end with XhoI site. Callus was induced on 2N6 media for 30 days and cultured for 36hrs on regenerated media"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Oryza sativa"
/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dev_stage="proliferated callus on 2N6 media for 30 days"
lab_host="E.coli SOLR"
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/db_xref="taxon:4530"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tissue_type="callus"
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root hair subtracted cDNA library
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RESULT 211
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EST.
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Contact: Kousaku Okubo, Naohiro Hori, Ryo Matoba,
Niiyama, Atsushi Fukushima, Yuko Kojima & Kenichi
Institute for Molecular and Cellular Biology
                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 17)

Okubo, K., Hori, N., Matoba, R., Nilyama, T., Fukushima, A., Kojima, Y., and Matsubara, K.
                                                                                                                                                                                                                                                     D11808
D11808.1 GI:2155083
                                                                                                                                                                                                                                                                                                    HUMHM01H11 Liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Gary Stacey
University of Missouri
108 Waters Hall, Columbia,
                                                                             Large scale cDNA sequencing for analysis of qualitative aspects of gene expression Nat. Genet. 2, 173-179 (1992)
                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                 mRNA sequence.
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Scheffler, B.E., Huang, S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 573-884-4752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2003)
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                                                  1345164
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fax: 573-882-0588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Expressed sequence tags from
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1. .17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                db_xref="taxon:3847"
/tissue_type="root hairs"
/clone_lib="Soybean root hair subtracted cDNA library
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'cultivar="Williams 82"
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                 Toshiyuki
Matsubara
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1 (bases 1 to 17)

Janssen, C.S., Barrett, M.P., Lawson, D., Quail, M.A., Harris, D.,

Bowman, S., Phillips, R.S. and Turner, C.M.

Gene discovery in Plasmodium chabaudi by genome survey sequencing

Mol. Biochem. Parasitol. 113 (2), 251-260 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AJ303755.1 GI:11140262
GSS; genome survey sequence.
Plasmodium chabaudi
Plasmodium chabaudi
                                                                                                                                                                                                                                                        Submitted (06-NOV-2000) Division of University of Glasgow, Joseph Black bases 39 to 55 (QL to SR).
                                                                                                                                                                                                                                                                                                                    2 (bases 1 to 17)
Janssen, C.S.
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1-3 Yamada-oka,S
                                                                                                                                                                                                                                                                                                         Direct Submission
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Liver HepG2 cell line."
/note="3'-directed regional cDNA library. Cleaved by MboI and transformed into E.coli."
                                                                                                                                                              /organism="Plasmodium chabaudi"
/mol_type="genomic DNA"
/db_xref="teaxon:5825"
/clone="PC7c5.plt"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="GDB:D0S8354E"
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/clone="hm01h11"
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PC7c5.plt,
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RESULT 213 PCH303755/c LOCUS

DEFINITION

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SOURCE ORGANISM

KEYWORDS

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REFERENCE

AUTHORS

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FEATURES COMMENT

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Percent Similarity:
Best Local Similarity:
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            Submitted (12-MAR-1999)
MIPS, Am Klopferspitz 1
                                                Duesterhoeft A., Lauber J.,
                                                                                                                                                                         Homo sapiens mRNA;
                                                                                           ) sapiens (human)
| ryota; Metazoa; Chordata; Craniata; Vertebrata;
|eria; Primates; Catarrhini; Hominidae; Homo.
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Submitted (06-NOV-2000) Division of
University of Glasgow, Joseph Black
bases 39 to 55 (QL to SR).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jakses 1 to 17)
Jakses, C.S., Barrett, M.P., Lawson, D., Quail, M.A., Harris, D.,
Bowman, S., Phillips, R.S. and Turner, C.M.
Gene discovery in Plasmodium chabaudi by genome survey sequencing
Mol. Biochem. Parasitol. 113 (2), 251-260 (2001)
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Plasmodium chabaudi
Plasmodium chabaudi
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
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                                                                                                                                                                                                                                                                                   standard; mRNA; EST; 18
                                                                                                                                                                                                 (Rel.
                                                                                                                                                sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Plasmodium chabaudi"
/mol type="genomic DNA"
/db_xref="taxon:5825"
/clone="PC7c5.plt"
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                                                                                                                                                                                                 Created)
Last upd
            9) to the EMBL/GenBank/DDBJ databases 18a D-82152 Martinsried, GERMANY
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e survey
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                                                                                                            Euteleostomi;
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Glasgow
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lasgow G12
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                                                                                                            Mammalia;
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Best Local Similarity:
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                                                         sequencing consultable

No sl sequence available at the RZPD in Berlin
This clone is available at the RZPD: Ressourcenzentrum, Ho
Please contact the RZPD: Ressourcenzentrum, Ho
Charlottenburg, GERMANY; Email: clone@
                                               Кеγ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Clone from S. Wiemann, sequenced by Qiagen within the cDNA sequencing consortium of the German Genome Project No s1 sequence available
This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 1 Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                                                                                           Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                        source
                                                                                                                                    Clone from
                                                                                                                                                                                              Blum H.,
                                                                                                                                                                                                                                                Eutheria;
                                                                                                                                                                                                                                                                                                                       Homo sapiens
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12-MAR-1999
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                                                                                                                                 S. Wiemann,
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                                                                                                                                                                                                                                                                                                                    mRNA;
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/db_xref="taxon:9606"
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/clone lib="434 (symonym: htes3). Vector pSport1; host
DH10B; sites Not1 + SalI"
/dev stage="adult"
/tissue_type="testis"
                                               Location/Qualifiers
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Last updated, Version
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@rzpd.de
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RESULT 214
HSM004368/c
ID HSM0043
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XX AL0398
XY I2-MAH
XY EST; 6
XY I1]
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RP 1-18
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RP 1-18
RP I1-18

Eutheria; Eukaryota; 12-MAR-1999 12-MAR-1999 AL039892.1 ᅜ Ş

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HSM004368

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AL039892;

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Query Match:

Score: Alignment

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                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
Hong/Eubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E.

Consortium (LLML) DNA Sequencing by: Barkeley MGC sequencing
project Clone distribution: MGC clone distribution information can
be found through the I.M.A.G.E. Consortium/LLML at:

www-bio.llnl.gov/bbrp/image/image.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center.
Trimming: cross match from University of Washingtion Genome Center
PHRAP suite. Poly-T Identification: patMatch.pl from Berkeley
Drosophila Genome Project. University of Washingtion Genome Center:
http://www.genome.washington.edu Low Quality Sequence: 18
contiguous PHRED high quality bases following vector sequence. Very
Low Quality Sequence: Trace file contained 18 contiguous distinct
peaks following vector sequence. Polyadenylation: Based upon the
presence of a XhoI site followed by a run of 14 or more I residues
and the beginning of the sequence, this cDNA insert was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scores:
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                                                                                                                                                                                                                                                  High quality sequence stop: 18.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                  polyadenylated.
Plate: LLCM7 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
Other_ESTs: 2821585.5prime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NIH-MGC http://mgc.nci.nih.gov/.
Mational Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/cell_line="MGC3"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_7"
/note="Organ: lung; Vector: pOTB7; Site_1: XhoI; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="DKPZp434B1823"
/clone_lib="434 (synonym: htes3). Vector pSport1; host
DH10B; sites Not1 + Sall"
/dev_stage="adult"
                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                           clone="IMAGE:2821585"
tissue_type="small cell carcinoma"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tissue_type="testis"
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COMMENT

FEATURES

quality sequence stop: 16.
Location/Qualifiers

/organism="Homo sapiens" /mol_type="mRNA"

/lab_noble_____/lab_noble__//la

/db xref="taxon:9606"
/clone="IMAGS:2821151"
/tissue type="small cell carcinoma"
/cell line="MGC3"
/cell line="MGC3"
/lab host="DH108 (phage-resistant)"
/clone lib="NHH MGC 7"

source

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REFERENCE
AUTHORS
TITLE
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ORGANISM
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AW250267
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VERSION
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Best Local Similarity:
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                                             Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LIML) DNA Sequencing by: Berkeley MGC sequencing project Clone distribution: MGC Clone distribution information can be found through the I.M.A.G.E. Consortium/LLML at: www-bio.llnl.gov/bbrp/1mage/image.html Base Calling / Quality Scores: PHRED from University of Washingtion Genome Center. Vector Trimming: cross match from University of Washingtion Genome Center PHRAP suite. Poly-T Identification: patMatch.pl from Berkeley Drosophila Genome Project. University of Washingtion Genome Center: http://www.genome.washington.edu_Low Quality Sequence: 16 contiguous PHRED high quality bases following vector sequence. Very Low Quality Sequence: Trace file contained 18 contiguous distinct
                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
Other ESTs: 2821151.3prime
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
peaks following vector sequence.
Plate: LLCM6 row: A column: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 18)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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AW250267.1 GI:6593260
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Score: rreq ORIGIN

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AW250449
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                           source
                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling

Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E.

Consortium (LLNL) DNA Sequencing by: Berkeley MGC sequencing

project Clone distribution: MGC clone distribution information can

be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html Base Calling / Quality

Scores: PHRED from University of Washingtion Genome Center:

Trimming: cross match from University of Washingtion Genome Center

Trimming: cross match from University of Washingtion Genome Center

PHRAP suite. PoTy-T Identification: patMatch.pl from Berkeley

Drosophila Genome Project. University of Washingtion Genome Center:

http://www.genome.washington.edu Low Quality Sequence: 18

contiguous PHRED high quality bases following vector sequence. Very

Low Quality Sequence: Trace file contained 18 contiguous distinct

peaks following vector sequence. Polyadenylation: Based upon the

presence of a XhoI site followed by a run of 14 or more T residues

nolyadenylared.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23 GlyArgArg 25
                                                                                                                                                                                                                                                                                                                                                     polyadenylated.

Plate: LLCM9 row: H column: 11
High quality sequence stop: 18.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 18)
NIH-MCC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1999)
Other_ESTs: 2822458.5prime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence.
/clone="IMAGE: 2822458"
/tlssue_type="small cell carcinoma"
/tlssue_type="small cell carcinoma"
/cell_line="McC3"
/lab_Tost="DH10B (phage-resistant)"
/clone_lib="NHH MGC_7"
/clone_Torgan: lung; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ ade by oligo-dT priming. Directionally
cloned into EcoRIXAhoI sites using the following 5'
adaptor: GGCACGAG(G). Size-selected >500bp for average
insert size_1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
                                                                                                                                                                                                                                                                                       organism="Homo sapiens"

mol_type="mRNA"
                                                                                                                                                                                                                                                                      db_xref="taxon:9606"
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US-09-966-880A-8 (1-198) x AW250449
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                    This clone is also available at Chinese National Human Genome Center at Shanghai, 351 Guo Shoujing Road, Zhangjiang Hi-Tech Pudong New Area, P.R.China. Please contact with Zhang Xu (xu.zhang@ion.ac.cn) or Han Zeguang (hanzg@chgc.sh.cn) PCR PRimers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Zhang Xu
Laboratory of Sensory Syst
Institute of Neuroscience
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DRABTE12 Rat DRG Library Rattus
                                                                                                                                                                                                                                                               Seq primer: POLYA=No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       320 Yue Yang Road, Shanghai 200031,
Tel: 86-21-64748700-121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: xu.zhang@ion.ac.cn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fax: 86-21-64713446
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/tissue_type="dorsal root ganglion"
/dev_stage="adult"
/clone_lib="Rat DRG Library"
                                                                                                                   /organism="Rattus norvegicus"
/mol_type="mRNA"
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/db_xref="taxon:10116"
                                                                                                                                                                                                                                       location/Qualifiers
                                                                        sex="male"
                                                                                                clone="DRABTE12"
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II RT (Life Technologies)."
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3.83e+06 3.00 100.00%

Matches: Conservative:

0 3 18

EST 06-NOV-2001

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RESULT 220
BG668047
                                                                                             Percent Similarity:
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Query Match:
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AUTHORS
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Laboratory of Sensory System
Institute of Neuroscience
320 Yue Yang Road, Shanghai 200031,
Tel: 86-21-64748700-121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Xiao,H.S., Huang,Q.H., Zhang,F.X., Bao,L., Lu,Y.J., Guo,C., Yang,L., Huang,W.J., Fu,G., Xu,S.H., Cheng,X.P., Yan,Q., Zhu,Z.D., Zhang,X., Chen,Z., Han,Z.G. and Zhang,X.

Identification of gene expression profile of dorsal root ganglion in the rat peripheral axotomy model of neuropathic pain proc. Natl. Acad. Sci. U.S.A. 99 (12), 8360-8366 (2002)
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DRABUA12 Rat DRG Library Rattus
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BG668047
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (xu.zhang@ion.ac.cn) or Han Zeguang (hanzg@chgc.sh.cn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: xu.zhang@ion.ac.cn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (bases 1 to 18)
                                                                                                                                                                                                                                                                                                                          /organism="Rattus norvegicus"
/mol type="mRNA"
/strain="Sprague-Dawley"
/db xref="taxon:10116"
/clone="DRABUA12"
                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                  /tissue_type="dorsal root ganglion"
/dev_stage="adult"
/clone_lib="Rat DRG Library"
                                                                                                                                                                                                                                                                                                             sex="male"
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Matches:
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                     REFERENCE
                                                                                         SOURCE
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Fax: 610-270-5598
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone_lib="HOA (Human Osteoarthritic Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
Directional"
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/mol_type="mRNA"
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Fax: 610-270-5598
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1 (bases 1 to 18)
                               Email: sanjay_kumar-1@gsk.com
Seg primer: T7
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Seq primer: T7.
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/clone lib="HOA (Human Osteoarthritic Cartilage)"
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Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen,M. and
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Directional"
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/clone_lib="HOA (Human Osteoarthritic Cartilage)"
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BG924473
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                                         BG925410.1
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Seq primer: T7.
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Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen
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/lab_host="E_coli_DH10_B"
/clone_lbe="NNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
Directional"
                                                                                                                                                                                                                                                                                                                                                                                       organism="Homo sapiens"
(mol_type="mRNA"
(db_xref="taxon:9606"
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BG925569
BG925569.1
                        GlaxoSmithKline
709 Swedeland Road, P.O. Tel: 610-270-7245
                                      UW2109
                                                Contact: Sanjay
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                                                                             21482651
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                                                  Kumar
           Box 1539, King
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Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and osteoarthritic cartilage cDNA libraries osteoarthr. Cartil. 9 (7), 641-653 (2001)
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1 (bases 1 to 18)
                                                                                                                        Kumar,S., Connor,J.R., Dodds,R.A., Halsey,W., Van Horn,M.,
Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen
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Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen,M.
Lark,M.W.
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Seq primer: T7.
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/lab_host="E.coli_DH10_B"
/clone_lib="HNC (Human Normal Cartilage)"
/note="Vector: psport I, Site_1: SalI, Site_2: NotI,
Directional"
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/db_xref="taxon:9606"
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Mismatches:
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Best Local Similarity:
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                Alignment Scores:
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BG925569
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18 bp mRNA linear EST 06-NOV-20 HNC5-1-E2.R HNC (Human Normal Cartilage) Homo sapiens cDNA, mRNA
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Sec primer: T7.
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Seq primer: T7.
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Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gowen,M. and
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Fax: 610-270-5598
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1 (bases 1 to 18)
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                                                              /tissue_type="cartilage"
/lab_host="E.coli DH10 B"
/clone_lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_
Directional"
                                                                                                                                           organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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/lab_host="E.coli DH10 B"
/clone_lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
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mol_type="mRNA"
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US-09-966-880A-8 (1-198) x BG927414 (1-18)
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HNC1-1-H3.R HNC (Human Normal
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Tel: 610-270-7245
Fax: 610-270-5598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GlaxoSmithKline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and osteoarthric cartilage CDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kumar, S., Connor, J.R., Dodds, R.A., Halsey, W., Van Horn, M., Mao, Sathe, G., Mui, P., Agarwal, P., Badger, A.M., Lee, J.C., Gowen, M. a
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 18)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: sanjay_kumar-1@gsk.com
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                                                                                                                                                                                                                                                                                                                                           /tissue_type="cartilage"
/lab host="E.coli DHIO B"
/clone_lib="HMC (Human Normal Cartilage)"
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Directional"
                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
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BG927414.1
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Tetrahymena thermophila
Tetrahymena thermophila
Tetrahymena thermophila
Ciliophora, Oligohymenophorea,
Eukaryota, Alveolata, Ciliophora, Tetrahymena.
University of Chicago
920 E. 58th Street, Chicago,
Tel: 773 702 4374
                                                                                              Frankel, J. and Klobutcher, L. EST from Tetrahvmena +harman
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                                             Contact: Turkewitz AP Molecular Genetics and Cell Biology
                                                                             Unpublished (2002)
                                                                                                                               Turkewitz,A.P.,
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                                                                                                                                                                                                                                                                             Tetrahymena thermophila cDNA, mRNA sequence.
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Seq primer: T7.
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Sathe, G., Mui, P., Agarwal, P., Badger, A.M., Lee, J.C., Gowen, M.
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                                                                                            from Tetrahymena thermophila, strain CU428.1, growing cells
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/lab_host="E.coli DH10 B"
/clone lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
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'db_xref="taxon:9606"
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                                                                                                                            Karrer, K.M., Jahn, C., Orias, E.,
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BM394638/c
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Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                             Molecular Genetics and Cell Biology University of Chicago
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Turkewitz, A.P., Karrer, K.M.,
Frankel, J. and Klobutcher, L.
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                                                                                                                                                                                                                     Seq primer: T3.
                                                                                                                                                                                                                                                                  Tel: 773 702 4374 Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                             Contact: Turkewitz AP
                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tetrahymena thermophila
Bukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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                                                                                                                                                                                                                                         Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                         920 E. 58th Street, Chicago,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tetrahymena thermophila
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/db_xref="taxon:5911"
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/note="Vector: BlueScript2 SK+; Details on library
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Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/note="Vector: BlueScript2 Sk., Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                                          /mol_type="mRNA"
/strain="CU428.1"
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Length:
Matches:
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Mismatches:
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                                                                                                                                      5009-0-3-F09.t.1 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence. BM397227
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                                           Tetrahymena thermophila
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Tetrahymena thermophila
Eukaryota, Alveolata, Ciliophora, Oligohymenophorea,
Hymenostomatida, Tetrahymenina, Tetrahymena.
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Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 606
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BM395123.1 GI:18195176
            1 (bases 1 to 18)
Turkewitz, A.P., Ka
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Frankel,J. and Klobutcher,L.
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/clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
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/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/mol_type="mRNA"
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|strain="CU428.1"
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Tetrahymena thermophila
Eukaryota; Alveolata; Clliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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Molecular Genetics and Cell Biology
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Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E., Frankel,J. and Klobutcher,L.
EST from Tetrahymena thermophila, strain CU428.1, growing
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Seq primer: T3.
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920 E. 58th Street, Chicago,
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Unpublished (2002)
Contact: Turkewitz AP
Molecular Genetics and Cell Biology
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920 E. 58th Street, Chicago,
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Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E.,
Frankel,J. and Klobutcher,L.
EST from Tetrahymena thermophila, strain CU428.1, growing cells
Unpublished (2002)
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                                                              BM398577 18 bp mRNA linear EST 17 5009-0-47-C10.t.1 Chilcoat/Turkewitz cDNA (large fraction)
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920 E. 58th Street, Chicago,
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18 bp mRNA linear EST 1
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                                  Tetrahymena thermophila BM398577
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Molecular Genetics and Cell Biology
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Hymenostomatida; Tetrahymenina; Tetrahymena.
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/strain="CU428.1"
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Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Eukaryota; Alveolata; Ciliophora; Tetrahymena.
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Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
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Unpublished (2002)
Contact: Turkewitz AP
Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 6063
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Turkewitz, A.P., Karrer, K.M.,
Frankel, J. and Klobutcher, L.
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                                                                                                             Tel: 773 702 4374
Fax: 773 702 3172
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BM401236
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Seq primer: T3.
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Contact: Turkewitz AP
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920 E. 58th Street, Chicago,
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Fax: 773 702 3172
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/db xref="taxon:5911"
/clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+, Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                              Location/Qualifiers
 /mol_type="mRNA"
/strain="CU428.1"
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Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., K. Frankel,J. and Klobutcher,L.
EST from Tetrahymena thermophila, strain CU428.1,
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Fax: 773 702 3172
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920 E. 58th Street, Chicago,
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Molecular Genetics and Cell Biology
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Hymenostomatida; Tetrahymenina; Tetrahymena.
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preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/mol type="mRNA"
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/note="Vector: BlueScript2 SK+; Details on library
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Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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                                                                                                                                                Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Amaranthaceae; Beta.

[ (bases 1 to 18)
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EST.
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E011887-024-004-L12-SP6 MPIZ-ADIS-024-inflorescence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOH602767971.R1 CSEQFXL35 adipose Sus scrofa cDNA, BM675715
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Adelson, D.L. and Gill, C.A.
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Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
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                                      Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
                                                                                                              Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M. 
Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Le
                                                                                                                                                                                                                                                      Beta vulgaris
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the inserts (AAGAATTCGATATCAAGCTTATCGATACCGTCGAGCTCGAG.
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/clone_Tib="CSEQFXL35 addipose;
/note="Organ; addipose tissue; Vector:
Site_1: NotI; Site_2: EcoRI; sequence
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/db_xref="taxon:9823"
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                                                                                                                                                                                                                                                 Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet gotton J. 32 (5), 845-857 (2002)
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Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,
Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
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ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
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Sp6-SalI-CCACGCTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note:
Sequencing granted in the context of the GABI-Beet
project, local PI: Dr. Katharina Schneider, coordinator:
Prof. Christian Jung; Sequence submission managed by
RZPD/GABI-Primary database: http://gabi.rzpd.de"
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/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not!;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not!;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not!;
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/note="Vector: pcmvs
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/clone="024-004-L12"
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllales; Amaranthaceae; Beta.
1 (Dases 1 to 18)
Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,
Drungowski, M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
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E011673-024-002-E13-SP6R MPIZ-ADIS-024-inflorescence
CDNA Clone 024-002-E13 5-PRIME, mRNA sequence.
                                                                       Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck Institute for Plant Breed
Carl-von-Linne Weg 10, 50829 Koeln,
Beta vulgaris
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primer: SP6r; ATTTAGGTGACACTATAGAAGA
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/note="Vector: pCMVSPORT6; Site 1: SalI; Site 2: NotI;
/note="Vector: pCMVSPORT6; Site 1: SalI; Site 2: NotI;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SP6-Sall-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RZPD/GABI-Primary database: http://gabi.rzpd.de"
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/mol_type="mRNA"
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/db_xref="taxon:161934"
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E012394-024-013-B09-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone
024-013-B09 5-PRIME, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Maranthaceae; Beta.

1 (bases 1 to 18)
                                                                                                               ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
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Location/Qualifiers
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/note="Vector: pcMVSPORT6; Site 1: Sali, Site 2: Not!,
/note="Vector: pcMVSPORT6; Site 1: Sali, Site 2: Not!,
/note="Vector: pcMVSPORT6; Site 1: Sali, Site 2: Not!,
/note="Vector: pcMVSPORT6; Site 2: Not!,
/note="Vector: pcMVSPORT6; Sali, Sa
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1. .16
/organism="Beta vulgaris"
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/lab_host="EMDH10B"
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US-09-966-880A-8 (1-198) x BQ586069 (1-18)
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Beta vulgaris
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S014468-024-013-P11-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone
024-013-P11 5-PRIME, mRNA sequence.
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Max-Planck-Institute for Plant Breeding Research
Carl-von Linne Weg 10, 50829 Koeln, Germany
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1 (bases 1 to 18)
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Location/Qualifiers
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/clone lib="MPIZ-ADIS-024-leaf"
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/note="Vector: pCMVSPORT6; Site 1: Sall; Sall; Site 2: NotI;
/note="Vector: pCMVSPORT6; Site 1: Sall; Sall;
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SP6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note:
Sequencing granted in the context of the GABI-Beet
project, local PI: Dr. Katharina Schneider, coordinator:
prof. Christian Jung; Sequence submission managed by
RZPD/GABI-Primary database:http://gabi.rzpd.de"
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/db_xref="taxon:161934"
/clone="024-013-B09"
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/cultivar="KWS2320 (double haploid, monogerm breeding
/organism="Beta vulgaris"
/mol_type="mRNA"
/cultivar="KWS2320 (double haploid,
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ORGANISM
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Plate: 13 row: P column: 11
Seq primer: SP6; CATACGATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                          Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
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[ (bases 1 to 18)
                                                                                                                                                                                                                                                                                                                         Carl-von-Linne Weg 10, 50829 Koeln, Fax: 00492215062851
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                                                                                                                                                                                                                                                                                               weisshaa@mpiz-koeln.mpg.de
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Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-Not!, primer sites and
/db_xref="taxon:161934"
/clone="024-013-P11"
/tissue_type="leaf"
/lab_host="EMDH10B"
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                                                                                                           line) "
                                                                                                                                                   Tom
                                                                                                                                                                                                                location/Qualifiers
                                                                                   db_xref="GABI:186486"
                                                                                                                                                                    organism="Beta vulgaris"
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/clone="024-013-P11"
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tivar="KWS2320 (double
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Score:

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/clone_lib="MPIZ-ADIS-024-leaf"
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not!;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not!;
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not!;
/note="Vector: provided by KWS
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-Not!, primer sites and
                                                                                                                                                                                                                                                                              Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by
                                                                                                                                                                                                                                                                                                                                                                    SPG-Sall-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Sequencing granted in the context of the GABI-Beet
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                                      3.83e+06
3.00
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Matches:
Conservative:
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BQ589347
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Query Match:
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ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and Radelof,U.

Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet get plant J. 32 (5), 845-857 (2002)
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                                                                                                                                                                                                                                                                                                             Email: weisshaa@mpiz-koeln.mpg.de
Insert Length: 18 Std Error: 0.00
Plate: 15 row: A column: 02
Seq primer: SP6; CATACGATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath, Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J.,
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/clone lib="MPIZ-ADIS-024-storage root" /note="Vector: pCMVSPORTS; Site 1: Sall; Site 2: N cDNA library from sugar beet, library provided by Kleinwanzlebener Saatzucht AG Einbeck, Germany, co
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Percent Similarity:
Best Local Similarity:
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E012616-024-017-G11-SP6 MPIZ-ADIS-024-storage root Beta vulgaris
CDNA clone 024-017-G11 5-PRIME, mRNA sequence.
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Plate: 17 row: G column: 11
Seq primer: SP6; CATACGATTTAGGTGACACTATAG
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
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[ (bases 1 to 18)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         weisshaa@mpiz-koeln.mpg.de
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/note="Vector: pcMVSPORT6; Site_1: SalI; Site_2: NotI;
/cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact
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SP6-SalI-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Sequencing granted in the context of the GABI-Beet
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                                                 orientation:
                                                                           b.schulz@kws.de; cloning sites SalI-NotI, primer sites
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/mol_type="mRNA"
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REFERENCE AUTHORS

COMMENT

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IEMBUG

JOURNAL MEDLINE TITLE ACCESSION VERSION

KEYWORDS

ORGANISM

DEFINITION

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RESULT 248

BQ591606

Query Match: DB:

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US-09-966-880A-8 (1-198) x BQ591606
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18 bp mRNA linear EST 06-DEC-2002
E012442-024-024-M20-SP6 MPIZ-ADIS-024-developing root Beta vulgaris
CDNA clone 024-024-M20 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Insert Length: 18 Std Error: 0.00
Plate: 24 row: M column: 20
Seq primer: SP6; CATACGATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet gaplant J. 32 (5), 845-857 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyliales; Amaranthaceae; Beta.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: weisshaa@mpiz-koeln.mpg.de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Max-Planck-Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Weisshaar B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Beta vulgaris
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                                                                                                                                                                /clone_lib="MPIZ-ADIS-024-developing root"
/note="Vector: pcMVSPORT6; Site 1: Sall; Site 2: Not1;
/note="Vector: pcMVSPORT6; Site 1: Sall; Site 2: Not1;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Binbeck, Germany, contact:
b.schulz@kws.de; cloning sites Sall-Not1, primer sites and
                                                    SP6-Sall-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet project, local FI: Dr. Katharina Schneider, coordinator:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             project, local PI: Dr. Katharina Schneider, coordir
Prof. Christian Jung; Sequence submission managed
RZPD/GABI-Primary database: http://gabi.rzpd.de"
Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de"
                                                                                                                                               orientation:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Beta vulgaris"
/mol_type="mRNA"
/cultivar="KWS2320 (double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                   /tissue_type="developing root"
/lab_host="EMDH10B"
                                                                                                                                                                                                                                                                                                                                                                          'clone="024-024-M20"
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BQ790001
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RESULT 251
C00629
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Eukaryota; mixed EST libraries.

1 (bases 1 to 18)

Asiegbu,F.O., Nahalkova,J. and Dean,R.A.
Selected Expressed sequence tags of cDNA clones from the
interaction of the root rot fungus (Heterobasidion annosum) with
seedling roots of Scots pine (Pinus sylvestris)
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hage005aH02 Heterobasidion annosum - Scots pine infection stage (HAGE) subtraction cDNA library Pinus sylvestris/Heterobasidion annosum cDNA clone hage005aH02, mRNA sequence.
                                                  თ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Fred O. Asiegbu
Dept. of Forest Mycology & Pathology
Swedish University of Agriculture, Box 7026,S-750 07,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: Fred.Asiegbu@mykopat.slu.
Seq primer: T7 primer.
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Fax: +46 18 30 92 45
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                                                                                  GluLeuLeu 60
                                                                                                                                                                                                                                                                                                       /dev_stage="Seedling roots of scots pine were infected for 6 days with H. annosum"
/clone lib="Heterobasidion annosum - Scots pine infection stage (HAGE) subtraction cDNA library"
/note="Vector: pT-Adv; Site_1: EcoRI; The subtractive hybridization cDNA library was constructed from scots pine roots infected for 6-days with mycelia of Heterobasidon annosum (FP5)."
                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="mRNA"
/db_xref="taxon:169015"
/clone="hage005aH02"
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Osaka University
1-3, Yamada-oka, Suita, Osaka Pref. 565, Japan
Tel: 06-877-5111(ex.3315)
Email: kousaku@imcb.osaka-u.ac.jp
We are not submitting the same cDNA sequence redundantly to DDBJ
                                                                                                   Okubo,K.

BodyMap; human gene expression database Unpublished (1995)

Contact: Okubo,K.

Contact: Okubo,K.
                                                                                                                                                                                                                                                                                                                 sequence.
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We are not submitting the same cDNA sequence redundantly to DDBJ
since 1993. For the abundance information of clones with this
sequence in this library and as well as in other 3'-directed
libraries, see 'http://www.imcb.osaka- u.ac.jp/bodymap'. The
sequences of the clones represented by this GS sequences is also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Osaka University
1-3,Yamada-oka, Suita, Osaka Pref. 565, Japan
Tel: 06-877-5111(ex.3315)
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BodyMap; human gene expression database Unpublished (1995)
Contact: Okubo,K.
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HUMGS0008172 Human adult
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 18)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                 Institute for Molecular and Cellular Biol
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/db_xref="taxon:9606"
/dev_stage="adult"
/clone_llb="Human adult (K.Okubo)"
/note="One or more human adult tissue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Homo sapiens"
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(K.Okubo)
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                                                                                                                                                                                                                                                                 1-3, Yamada-oka, Suita, Osaka Pref. 565, Japan Tel: 06-877-5111(ex.3315)
Email: kousakusimcb.osaka-u.ac.jp
We are not submitting the same cDNA sequence redundantly to DDBJ with this since 1993. For the abundance information of clones with this sequence in this library and as well as in other 3'-directed libraries, see 'http://www.imcb.osaka- u.ac.jp/bodymap'. The sequences of the clones represented by this GS sequences is also
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                                                                                                                                                                                                                                                                                                                                                                                               Osaka University
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Unpublished (1995)
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Mammalia, Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="mRNA"
/db_xref="taxon:9606"
/dev_stage="adult"
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/note="One or more human adult tissue"
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/clone_lib="Human adult (K.Okubo)"
/note="One or more human adult tissue"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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CA851280.1
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D12A08_B20_02.ab1
cDNA clone D12A08
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Email: kousaku@imcb.osaka-u.ac.jp

We are not submitting the same cDNA sequence redundantly to DDBJ

since 1993. For the abundance information of clones with this

sequence in this library and as well as in other 3'-directed

libraries, see ' http://www.imcb.osaka- u.ac.jp/bodymap'. The

sequences of the clones represented by this GS sequences is also
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Okubo r
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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                                                         Glycine max
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/mol_type="mRNA"
/db_xref="raxon:9606"
/dev_stage="adult"
/clone_lib="Human adult (K.Okubo)"
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(K.Okubo) Homo sapiens cDNA 3', mRNA
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                                                                                                                           Soybean Genomics and Improvement Laboratory (SGIL) US Department of Agriculture (USDA), ARS, PSI Bldg.006, Rm 118, 10300 Baltimore Ave., Beltsville
                                                                                                                                                                                                  Alkharouf, N.W., Khan, R. and Matthews, B.F. Analysis of expressed sequence tags from infected by the soybean cyst nematode Unpublished (2002)
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Glycine max (soybean)
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Alkharouf,N.W., Khan,R. and Matthews,B.F.
Analysis of expressed sequence tags from roots of resistant soybean infected by the soybean cyst nematode
Unpublished (2002)
Contact: Alkharouf, N.W.
                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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CA851607.1 GI:33388400
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US Department of Agriculture (USDA), ARS, PSI
Bldg.006, Rm 118, 10300 Baltimore Ave., Beltsville, MD 20705-2350,
                                                                                                                                                                                 Contact: Alkharouf, N.W.
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Fax: 301 504 5728
Email: alkharon@ba.ars.usda.gov.
Location/Qualifiers
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                               301 504 5750
301 504 5728
1: alkharon@ba.ars.usda.gov.
Location/Qualifiers
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/dev_stage="Seedlings"
/clone_lib="cDNA peking library 2, 4 day SCN3"
/clone_lib="cDNA peking library 5, 4 day SCN3"
/note="Vector: pBluescript SK-; cDNA clones from mRNA
extracted from Peking roots 2 and 4 days past invasion."
organism="Glycine max"
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/clone="D12A08"
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/cultivar="Peking"
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CD486685/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18 bp mRNA linear EST 01-JUL-2003 CRH5.3A10 Cotton Root and Hypocotyl Lambda ZIPLOX Library (CRH) Gossypium hirsutum cDNA clone CRH5.3A10 5', mRNA sequence. CD486685 CD486685.1 GI:31407650 EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Different Gene Expression Responses in Cotton Root and Hypocotyl tissues during infection with Fusarium Wilt Disease Unpublished (2003) Contact: Caitriona Dowd, Helen McFadden Commonwealth Scientific and Industrial Research Organisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Black Mountain, Canberra, ACT,
Tel: 61 2 6246 4914, 6246 5377
Fax: 61 2 6246 5000
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Black Mountain, Canberra, ACT
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Gossypium hirsutum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: Caitriona.Dowd@csiro.au, Helen.McFadden@csiro.au
Vector clipped sequences Bases 1-17 (GTCGACCCACGCGTCCG): SalI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Division of Plant Industry
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Dowd, C., Wilson, I. and McFadden, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer: M13 reverse primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   quality sequence stop: :
Location/Qualifiers
/note="Vector: Lambda ZIPLOX; Site 1: SalI; Site 2: NotI; mRNA was prepared from root and hypocotyl tissues of the cotton cultivar DeltaRMERALD. cDNA was synthesised from a NotI-oligodT primer/adapter using the manufacturers protocols (Life Technologies) and then ligated to a SalI adapter to facilitate directional cloning. The cDNA was cloned into the SalI and NotI sites of the Lambda ZIPLOX
                                                                                                                                                                                /tissue_type="Root and hypocotyl tissues"
/dev_stage="5 day old seedlings"
/lab_host="y1090(ZL)"
/clone_lib="Cotton Root and Hypocotyl Lambda ZIPLOX
Library_(CRH)"
                                                                                                                                                                                                                                                                                                                                 /mol_type="mRNA"
/cultivar="DeltaEMERALD"
/db_xref="taxon:3635"
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/dev_stage="Seedlings"
/clone lib="cDNA Peking library 2, 4 day SCN3"
/clone lib="CDNA peking library 2, 4 day SCN3"
/note="Vector: pBluescript SK-; cDNA clones from mRNA
/note="Vector Peking roots 2 and 4 days past invasion."
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/db_xref="taxon:3847"
/clone="D15F01"
                                                                                                                                                                                                                                                                                                                  clone="CRH5.3A10"
                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Gossypium hirsutum"
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T, 2601, Australia
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RESULT 258
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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CF308804.1 GI:33680565
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ABF--02-M04.gl ABF3-overexpressing transgenic rice plasmid cDNA
library (ABF) Oryza sativa cDNA clone ABF--02-M04, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oryza sativa
Oryza sativa
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Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 82 31 330 6193 Fax: 82 31 321 6355
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    LeuArgAsn
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Location/Qualifiers
1. .18
                                                                                                                                                                                                            /tissue_type="leaf"
/dev_stage="l4 days after germination"
/lab_host=="E.coli DH10B"
/clone_lib="R.coli DH10B"
/clone_lib="ABF3-overexpressing transgenic rice plasmid cDNA library (ABF)"
/note="Vector: pCR4-TOPO; Site 1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             phage vector (Life Technologies). Constructed by Caitriona
Dowd and Helen McFadden."
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clone="ABF--02-M04"
                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/cultivar="Nackdong"
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Oryza sativa
                                               Oryza sativa
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Kim, J. S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                  Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech Inc., Division of Bioscience and Bioinformatics, MyongJi University
Yongin, Kyeenggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                Oryza sativa
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza
                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 bp mRNA linear EST 15-AUG-2003 HD--06-N14.bl OsHDACl-overexpressing transgenic rice plasmid cDNA library (HD) Oryza sativa cDNA clone HD--06-N14, mRNA sequence.
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Genomics and Genetics Institute, GreenGene Biotech Inc.,
of Bioscience and Bioinformatics, MyongJi University
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
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CF317226.1 GI:33688987
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               bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="E.coli DH10B"
/clone lib="DSHDAC1-overexpressing transgenic rice plasmid
cDNA library (HD)"
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/cultivar="Nackdong"
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VERSION
KEYWORDS
SOURCE
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CF319738/c
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AUTHORS
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Best Local Similari
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 bp mRNA linear EST 15-AUG-2003
HD--10-F16.bl OsHDACl-overexpressing transgenic rice plasmid cDNA
library (HD) Oryza sativa cDNA clone HD--10-F16, mRNA sequence.
CF319738
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., L.,
Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoldeae; Oryzeae; Oryza.
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                                                                                                                                                                                                                                                                                                                                                                                 Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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                                                                                                                                                                                                                                                                                                                                                               Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Nahm B.H.
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reverse
derived
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(note="Vector: pCR4-TOPO; Site 1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression
  /note="Vector: pCR4-TOPO; Site_1: EcoRI; Callus was treated with ABA(20um) for lnr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression
                                                              /clone lib="OsHDAC1
cDNA library (HD)"
/note="Vector: pCR4
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/dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="E.coli_DH10B"
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                                                                                                            /dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="E.coli DH10B"
/clone_lib="OsHDAC1-overexpressing transgenic rice plasmid
                                                                                                                                                                                                                                                /mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                             tissue_type="callus"
                                                                                                                                                                                                      clone="HD--10-F16"
                                                                                                                                                                                                                         db_xref="taxon:4530"
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RESULT 263
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HDN--02-N01.g1 OsHDAC1-overexpressing transgenic rice lambor cDNA library II (HDN) Oryza sativa cDNA clone HDN--02-N01,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University

Yongin, Kyeenggi, Korea

Tel: 82 31 330 6193

Fax: 82 31 321 6355
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Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       bhnahm@ggbio.com, bhnahm@bio.myongji.ac.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                               /organism="Oryza sativa"
/mol_type="mRNA"
/cultivar="Nackdong"
/db xref="taxon:4530"
/clone="HDN--02-N01"
                                                                                                                                                                                                                                                                               /tissue_type="callus"
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/lab_host="E.coli SOLR"
/clone_lib="OsHDAC1-overexpressing transgenic rice lambda
phage_CDNA_library_II (HDN)"
phage_CDNA_library_II (HDN)"
                                                                                                                                                                                                                                      /note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: XhoI; cDNA was inserted into lambda Uni-ZAP XR vector at 5' end with EcoRI and 3' end with XhoI site. mRNA was
                                                                                                                                                                                                             derived from rice Histone Deacetylase overexpression line."
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Mismatches:
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CF330870/c
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Query Match:
DB:
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Oryza sativa
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., I
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
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Ehrhartoideae; Oryzeae; Oryze.

1 (bases 1 to 18)

1 (bases 1 to 18)

Song, S. I., Kim, J. Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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JMT--01-A23.g1 AtJMT-overexpressing
library (JMT) Oryza sativa cDNA clor
CP332520 GI:33813259
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
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CF330870
                                         Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, Ehrhartoideae, Oryzeae, Oryza.

1 (bases 1 to 18)
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/mol_type="mbNA"
/cultivar="Mackdong"
/cultivar="Mackdong"
/db xref="taxon:4530"
/db xref="taxon:4530"
/clone="NACL--06-M07"
/tissue_type="callus"
/tis
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Location/Qualifiers
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1 AtJMT-overexpressing transgenic rice
Oryza sativa cDNA clone JMT--01-A23, m
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Lee, T.H., Shin, Y.C.,
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AUTHORS
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VERSION
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CF333354
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                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Shrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 18)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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Unpublished (2003)
Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
Yongin, Kyeenggi, Korea
Tel: 82 31 330 6193
Tea: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                              CF333354 IS D MRNA linear BST 18-AUG-2003 JMT--02-D13.gl AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT) Oryza sativa cDNA clone JMT--02-D13, mRNA sequence. CF333354 GI:33814976
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                                                                                                                                                                                              of Bioscience and Bioinformatics,
                                                                                                                                                                                                               Genomics and Genetics Institute, GreenGene Biotech Inc.; Division
                                                                                                                                                                                                                                   Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                             Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
Location/Qualifiers
                                                                                                                                         in, Kyeonggi, Ko
82 31 330 6193
82 31 321 6355
                                                                                                        bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
Location/Qualifiers
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/lab_host="E.coli DH10B"
/clone lib="AtUMT-overexpressing transgenic rice plasmid cDNA library (JMT)"
/nott="Vector: pCR4-TOPO; Site_1: EcoRI; Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was pepared from Arabidopsis Jasmoinate Carboxyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:4530"
/clone="JMT--01-A23"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             methyltransferase overexpression line.
 /mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:4530"
/clone="JMT--02-D13"
                                                                     organism="Oryza sativa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tissue_type="leaf"
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100.00%
                                  3.83e+06
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US-09-966-880A-8 (1-198) x CF333354 (1-18)
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Genomics and Genetics Institute, GreenGene Biotech Inc.,
of Bioscience and Bioinformatics, Myongdi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 18)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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JMT--03-M11.g1 AtJMT-overexpressing transgenic rice plasmid cDNA
library (JMT) Oryza sativa cDNA clone JMT--03-M11, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                           Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
    Location/Qualifiers
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/dev stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone_lib="AtdWT-overexpressing transgenic rice plasmid cDNA_library (JWT)"
                          /note="Vector: pCR4-TOPO; Site_1: EcoRI; Oligo-capped mRNA
was reverse transcribed and then used for PCR. mRNA was
pepared from Arabidopsis Jasmoinate Carboxy1
                                                                                                      /organism="Oryza sativa"
/mol type="mRNA"
/culTivar="Nackdong"
/culTivar="Nackdong"
/db xref="taxon:4530"
/clone="JMT--03-M1"
/tissue_type="leaf"
/dev stage="14 days after germination"
/lab host="E.coli DH10B"
/clone_lib="AtJMT-overexpressing transgenic rice plasmid
cDNA_library (JMT)"
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pepared from Arabidopsis Jasmoinate Carboxyl
methyltransferase overexpression line."
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Conservative: Mismatches:

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Schistosoma mansoni
Eukaryota; Metazoa; Platyhelminthes; Trematoda; Digenea;
Strigeidida; Schistosomatoidea; Schistosomatidae; Schistosoma.
                                                                                        SCMRAP0216
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HUM000C318 Liver HepG2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Large scale cDNA sequencing for analysis of quantitative and qualitative aspects of gene expression Nat. Genet. 2, 173-179 (1992)
                                                        L76122.1 GI:1196860
                                                                                                     L76122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Kousaku Okubo, Naol
Niiyama, Atsushi Fukushima,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Okubo, K., Hori, N., Matoba, R., Niiyama, T., and Matsubara, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           D11637.1 GI:2148229
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D11637
                                  Schistosoma mansoni
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Location/Qualifiers
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                                                                            P0216 G2/KS adult
SMRAP0216, mRNA
                                                                                                                                                                                                                                                                                                                       /clone_lib="Liver HepG2 cell line."
/note="3'-directed regional cDNA library. Cleaved by and transformed into E.coli."
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                                                                                                                                                                                                                                                                                                                                                                                                                organism="Homo sapiens"
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db_xref="GDB:D087522E"
db_xref="taxon:9606"
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                                                                                        adult worm mini-library Schistosoma mansoni cDNA
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cell line.
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ima, Yuko Kojima
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ma & Kenichi
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Matsubara
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Berlin-Charlottenburg,
                                                                   Clone from S. Wiemann, se
sequencing consortium of
                                                                                                                                          Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                               Blum
                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, H
                                                  No si sequence available
                                                                                                                           MIPS,
                                                                                                                                                                                                                                                                                         Homo sapiens (human)
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12-MAR-1999
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                                                                                                                                                                                                                                                                                                                           expressed sequence tag.
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                                                                                                                           Am Klopferspitz 18a D-82152 Martinsried,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 18)
Neto,E.D., Harrop,R, Correa-Oliveira,R, Wilson,R.A., Pena,S.D. and Simpson,A.J.G.
Minilibraries constructed from cDNA generated by arbitrarily primed RT-PCR: an alternative to normalized libraries for the generation of ESTs from nanogram quantities of mRNA
Gene 186 (1), 135-142 (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Neto,E.D., Harrop,R.,
Pena,S.D. and Simpson,A.J.
Location/Qualifiers
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                                                                                                                                                                               Bauersachs S., Mewes W., Gassenhuber J., Wiemann
                                                                                                                                                                                                                                                                                                                                                                                                (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                             mRNA; EST DKFZp434C1822_r1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="SMRAP0216"
/clone_lib="G2/KS adult worm mini-library"
/clone_lib="G2/KS adult worm mini-library"
/note="A mini-library was made by cloning products deriv from RNA-arbitrarily primed PCR (RAP PCR) profiles into the pUC 18 vector: Reverse transcription of adult worm mRNA was primed with G2and subsequent PCR amplification was performed in the presence of primer KS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       db_xref="taxon:6183"
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GERMANY; Email:
                                                                     sequenced by of the German
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Matches:
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                                                                     LMU within the cDNA Genome Project
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clone@rzpd.de
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                Heubnerweg
                                                                                                                           GERMANY
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Location/Qualifiers

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HSM007596
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    Seguence 19
                                                                                                                                                                                                                                                                                                            This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, HG Berlin-Charlottenburg, GERMANY; Email: clone@
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                                                                                                                                                                                                                                                                                                                                                                                                        sequencing consortium of
                                                                                                                                                                                                                                                                                                                                                                                                                               Clone from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Blum H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eutheria;
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12-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                        S. Wiemann,
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    8만;
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/mol_type="mRNA"
/organism="homo sapiens"
/clone="DKEZP434C1822"
/clone lib="434 (synonym: htes3). Vector pSport1; host DH10B; sites NotI + SalI"
/dev_stage="adult"
                                                                /organism="Homo sapiens"
/clone="DKRZp434C1822"
/clone_lib="434 (symonym: htes3). Vector pSport1; host
DH10B; sites NotI + SalI"
/dev_stage="adult"
                                                                                                                                                                                                                                                                   Location/Qualifiers
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/mol_type="mRNA"
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                                           /tissue_type="testis"
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  C; 6 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mewes W., Gassenhuber J., Wiemann S.;
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clone@rzpd.de
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KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19 bp mRNA linear EST 04-am21b11.sl Soares NFL T GBC S1 Homo sapiens cDNA clone IVAGE:1467453 3' similar to TR:093040 Q93040 TIFIBETA ZINC PROTEIN. [1] ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15 TTCCGGACC 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seq primer: -40m13 fwd. ET from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Trace considered overall poor quality
Possible reversed clone: similarity on wrong
Possible reversed clone: polyT not found
Insert Length: 1216 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tumor Gene Index Unpublished (1997)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
Mational Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                     /clone lib="Soares NFL T GBC S1"
/clone lib="Soares NFL T GBC S1"
/note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with
a modified polylinker; Site 1: Not I; Site 2: Eco RI;
Equal amounts of plasmid DNA from three normalized
libraries (fetal lung NbHL19W, testis NHT, and B-cell
NCI_CGAP_GCB1) were mixed, and ss circles were made in
vitro. Following HAP purification, this DNA was used as
tracer in a subtractive hybridization reaction. The driver
was PCR-amplified cDNAs from pools of 5,000 clones made
from the same 3 libraries. The pools consisted of
I.M.A.G.E. clones 297480-322087, 682632-687239,
726408-728711, and 729096-731399. Subtraction by Bento
Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:1467453"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Homo sapiens"
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Indels:
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Euteleostomi;
; Homo.

EST 04-JAN-1999

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RESULT 273
AA903030
LOCUS
 ACCESSION
VERSION
KEYWORDS
                                                                                                   RESULT 274
AA918795
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19 bp mRNA linear EST 10-JUN-1998 ol69c05.sl NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:1534856 3' similar to TR:Q39599 EXTENSIN.; contains TAR1.b2 TAR1 repetitive element;, mRNA sequence.
AA918795.1 GI:3058685
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unknown library type
Trace considered overall poor quality
Insert Length: 714 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
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1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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AA903030.1 GI:3038153
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                                                                                                                                                                                                                                                                                                                                                               /tissue_type="leiomyosarcoma"
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/db_xref="taxon:9606"
/clone="IMAGE:1517462"
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Matches:
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AA918795/c
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Best Local Similarity:
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Unpublished (1997)
                                                                                    Homo sapiens (human)
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1.52%
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19 bp mRNA linear EST 10-JUN-1998 ol69c05.81 NCI CGAP Kid3 Homo sapiens cDNA clone IMAGE:1534856 3' similar to TR:\(\overline{Q}\)39599 EXTENSIN. ;contains TAR1.b2 TAR1 repetitive element ;, mRNA sequence.
AA918795
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1 (bases 1 to 19)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 19)
NCI-CGAP http://www.ncbi.nlm.nih.gov/nciogap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP);
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//lab host="DHIOB"
//clone=lib="NCI CGAP Kid3"
//clone=lib="NCI CGAP Kid3"
//clone="Organ: kidney; Vector: pT7T3D-Pac (Pharmacia) with note="Organ: kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; lst strand cDNA was primed with a Not I - oligo(dT) primer, double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. mRNA source: 2 pooled kidneys. Library went through one round of normalization. Library constructed by Bento Soares and M. Fatima Bonaldo. "
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/db_xref="taxon:9606"
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                                                                                                                                                       Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 19)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP);
                                                                                                                                                                                                                                                                                                                                                                                                                  AA932041 19 bp mRNA linear EST 07-JUL co35h05.s1 NCI CGAP Lu5 Homo sapiens cDNA clone IMAGE:1568217 similar to SW:NU4M_PANTR P03906 NADH-UBIQUINONE OXIDOREDUCTASE
                     Contact: Robert Strausberg, Ph.D.
Email: cgapbs-Yemail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
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Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
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Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                   Unpublished (1997)
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                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                   AA932041.1 GI:3087083
                                                                                                                                                                                                                                                                                                                                                                                     CHAIN 4 ;, mRNA sequence.
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Library Preparation: M. Bento
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="NCI_CGAP_Kid3"
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Soares,
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                                      Williams, S.A., Lizotte-Waniewski, M., Laney, S., Wenhong, L., Hillier, L., Allen, M., Bowles, L., Geisel, S., Jost, S., Kucab Martin, J., Steptoe, M., Theising, B., White, Y., Wylle, T., Chappell, J., Person, B., Gibbons, M., Harvey, N., Pape, D., Chamberlain, A., Morales, R., Schurk, R., Ritter, E., Kohn, S., Underwood, K. and Marra, M. Molecular Parasicology CvL3
                                                                                                                                                                                                                                                                                                                                                                                                                       AA934303 19 bp mRNA linear EST 26-MAR-1999 SWOVL3CAN12H12 Onchocerca volvulus infective larva cDNA (SAW94WL-OvL3) Onchocerca volvulus cDNA clone onch672 5' similar to TR:033571 Q33571 ATPASE SUBUNIT 6 ;, mRNA sequence.
Unpublished (1998)
Contact: Steven A. Williams
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Trace considered overall poor quality Insert Length: 525 Std Error: 0.00 Seq primer: -40m13 fwd. ET from Amersham High quality sequence stop: 1.
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1 (bases 1 to 19)
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DNA Sequencing by: Washington University Genome
Clone distribution; NCI-CGAP clone distribution
found through the I.M.A.G. E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Nematoda; Chromadorea; Spirurida; Filarioidea;
                                                                                                                                                                                                                                                                                                                          Onchocerca volvulus
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db_xref="taxon:9606"
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                                                                                                                                                                          Kucaba, T.,
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Smith College Department of Biological Sciences
Department of Biological Sciences, Clark Science Center, Smith
College, Northampton, MA, 01063, USA
Tel: 4135853826
Fax: 4135853786

Molecular Parasitology

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REFERENCE
AUTHORS
TITLE
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AI016864/c
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ou27c11.x1 Soares NFL T GBC_S1 Homo sapiens cDNA clone IMAGE:1627508 3' similar to TR:Q35989 Q35989 CYTOCHROME C OXIDASE SUBUNIT 1 ; mRNA sequence.
                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 19)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP);
                                                                                                                                                                                                                                                  Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SWOVL3CAN12H12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Vector: lambda UniZap XR; Site 1: EcoR I; Site 2: Xho I; Cutaneous filarial nematode parasite of humans. mRNA was prepared from third stage infective larvae of Onchocerca volvulus isolated from mosquitoes 10 days after infection and converted to double stranded cDNA using reverse transcriptase and oligo(dT) followed by RNase H and DNApol I. The library had 1.8 x 10E5 independent recombinants and average insert size was 900 base pairs. The library was constructed by Wenhong Lu. The library is available from Dr. S.A. Williams, email genome@smith.edu."
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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/clone_lib="Onchocerca volvulus infective larva cDNA
(SAW94WL-OvL3)"
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/clone="onch672"
                                                                                  Socation/Qualifiers
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AI017940
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ORGANISM
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Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Trace considered overall poor quality
Insert Length: 1853  Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D. Email: cgapbs-romail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 19)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/organism="Homo sapiens"
/mol_type="meNAA"
/mb_xref="taxon:9606"
/clone="IMAGE:1627183"
/clone="IMAGE:1627183"
/clone="IMAGE:1627183"
/clone=lib="Soares NFL_T GBC SI"
/clone=lib="Soares NFL_T GBC SI
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//Clone lib="Soares NFL T_GBC_S1"
//note="Organ: pooled; Vector: p7773D-Pac (Pharmacia) with
a modified polylinker; Site 1: Not I; Site_2: Eco RI;
Equal amounts of plasmid DNA from three normalized
libraries (fetal lung NbHL19W, testis NHT, and B-cell
NCI_CGAP_GCB1) were mixed, and ss circles were made in
vitro. Following HAP purification, this DNA was used as
tracer in a subtractive hybridization reaction. The driver
was PCR-amplified cDNAs from pools of 5,000 clones made
from the same 3 libraries. The pools consisted of
I.M.A.G.E. clones 297480-302087, 682632-687239,
726408-728711, and 728096-731399. Subtraction by Bento
Soares and M. Fatima Bonaldo. "
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/lab_host="DH10B"
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US-09-966-880A-8 (1-198) x AI017940 (1-19)
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This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Trace considered overall poor quality
Insert Length: 897 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
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1 (Dases 1 to 19)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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AI077581.1 GI:3411989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1997)
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                                                                                                                                                                                                                                                                                               /tissue_type="senescent fibroblast"
/lab_host="DH10B (ampicillin resistant)"
/clome_lib="Soares_senescent_fibroblasts_NbHSF"
/clome_tib="Soares_senescent_fibroblasts_NbHSF"
/note="Vector: p7773D (Pharmacia) with a modified
polylinker V_TYPE: phagemid; Site_1: Not I; Site_2: Eco
RI; lst_strand_cDNA was primed with a Not I - oligo(dT)
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from the same 3 libraries. The pools consisted of
I.M.A.G.E. clones 297480-302087, 682632-687239,
726408-728711, and 729096-731399. Subtraction by Bento
Soares and M. Fatima Bonaldo. "
                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                   clone="IMAGE:1666926"
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AI078728
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                   σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 575 Std Error: 0.00
Seq primer: -40nl3 fwd. ET from Amersham
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A1079728 19 bp mRNA linear EST 24-S 0912f07.81 Soares senescent fibroblaste NbHSF Homo saptens clone IMAGE:1665637 3' similar to SW:POR2 HUMAN P45880 VOLTAGE-DEPENDENT ANION-SELECTIVE CHANNEL PROTEIN 2; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 19)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumor Gene Index
Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                        /tissue_type="senescent fibroblast"
/lab_host="PH10B (ampicillin resistant)"
/clone_lib="Soares senescent_fibroblasts_MbHSF"
/clone_tib="Soares senescent_fibroblasts_MbHSF"
/note="Vector: pT7T3D (Pharmacia) with a modified
polylinker V_TYPE: phagemid; Site_1: Not I; Site_2: Eco
RI; lst strand cDNA was primed with a Not I - oligo(dT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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REFERENCE AUTHORS TITLE

COMMENT

ORIGIN

Pred. No.: Alignment

Percent Similarity:

SOURCE ORGANISM

Homo sapiens

VERSION

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DEFINITION

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118 Аврагдіув 120

1 GACCGCAAA 9

Query Match: DB:

Percent Similarity: Best Local Similarity:

Score: Pred. No.: Alignment

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AI147066/c
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AI155325
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                                           AI155325 19 bp mRNA linear EST 30-SEP-1998 ud88a05.r1 Soares NMPu Mus musculus cDNA clone IMAGE:1477904 5' similar to TR:Q62084 Q62084 PHOSPHOLIPASE C NEIGHBORING ;, mRNA
    sequence.
AI155325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Seq primer: -40ml3 fwd. ET from Amersham
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 19)
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Unpublished (1997)
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                                                                                                                                                                               AGGITGICA 4
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone lib="Source NSF P8 9W OT PA P S1"
/note="Organ: pooled; Vector: pTTT3D-Pac (Pharmacia) with
/note="Organ: pooled; Vector: pTTT3D-Pac (Pharmacia) with
a modified polylinker; Site_1: Not I; Site_2: Eco RI;
Equal amounts of plasmid DNA from five normalized
libraries were mixed, and ss circles were made in vitro.
Following HAP purification, this DNA was used as tracer in
a subtractive hybridization reaction. The driver was
PCR-amplified cDNAs from pools of 5,000 clones made from
the same 5 libraries. The pools consisted of the following
libraries and cloneIDs: Soares NbHSF pool 1:
309384-310919, 323208-325895 Soares Nb2HP pool 1:
145032-147335, 147720-148103, 148872-149255, 15002 -
150407, 151176-152327 Soares NbHSP pool 1:
758280-760583, 772104-774407 Soares NbHPA pool 1:
304776-306311, 320136-322823, 326280-32663 Soares NbHOT
pool 1: 723720-726407, 739080-740999 Subtraction by Bento
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="IMAGE:1509591"
/lab_host="DH10B"
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sapiens (human)
sapiens
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US-09-966-880A-8 (1-198) x AI155325 (1-19)
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WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Trace considered overall poor quality Possible reversed clone: similarity on wrong Seq primer: -28ml3 revz ET from Amersham High quality sequence stop: 1.

Location/Qualifiers
                                                                                 AI360784 19 bp mRNA linear EST 15-FEB-1999 qx98g07.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2010588 3's similar to TR:Q41707 Q41707 EXTENSIN CLASS 1 PROTEIN PRECURSOR.;
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Mammalia; Eutberia; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus (house mouse)
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AI360784.1 GI:4112405
                                                      mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: mouseest@watson.wustl.edu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone lib="Soares_NMPu"
/clone lib="Soares_NMPu"
/note="Organ: uterus; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from pregnant mouse uterus, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized, Library was constructed by Bento Soares and M. Fatima Bonaldo. "
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/lab_host="DH10B"
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/mol_type="mRNA"
/db_xref="taxon:10090"
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Mismatches:
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1 (bases 1 to 19)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
                                                                                                                                                                                                            ta07909.x1 NCI CGAP Brn23 Homo sapiens cDNA clone IMAGE:2043424 3' similar to TR:\(\overline{Q}26195\) Q26195 PVA1 GENE. ; contains L1.b3 L1 repetitive element ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 973 Std Error: 0.00
Seq primer: -40UP from Gibco
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1 (bases 1 to 19)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurole
Disorders and Stroke, Brain Tumor Genome Anatomy Project
                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                  Homo sapiens
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                                                                                                                                                   sapiens (human)
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(mol type="mRNA"

/db xref="taxon:9606"

/clone="IMAGE:2010588"

/clone="IMAGE:2010588"

/tissue type="glioblastoma (pooled)"

/lab_host="DH108"
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                                                                                                                                                                                                                                                                 A1431460 RST 13-APR-1999 THANA Linear EST 13-APR-1999 THA0601.x1 NCI CGAP Lym12 Homo sapiens cDNA clone IMAGE:2120736 3' similar to TR:004117 Q04117 SALIVARY PROLINE-RICH PROTEIN RP4 PRECURSOR. ; contains element MSR1 repetitive element ; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Trace considered overall poor quality Insert Length: 536 Std Error: 0.00 Seq primer: -40UP from Gibco High quality sequence stop: 1.
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Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
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                                 Tumor Gene Index
Unpublished (1997)
                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 19)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D., Myrna R.
Contact: Robert Strausberg,
Email: cgapbs-r@mail.nih.go
                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:2043424"
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/lab_host="DH10B"
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AI433480
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Best Local Similarity:
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLML at:
www-bio.llnl.gov/bbrp/image/image.html
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                                                                                                      Trace considered overall poor quality Insert Length: 1003 Std Error: 0.00
                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                              Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                           Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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_primer: -40UP from Gibco
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primer: -40UP from Gibco
h quality sequence stop: 1.
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                                      quality sequence stop: 1.
Location/Qualifiers
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/note="Organ: lymph node, Vector: pCMV-SPORT6; Site_1:
SalI; Site_2: NotI; Cloned unidirectionally. Primer:
Oligo dT. Average insert size 1.25 kb. Life Technologies
catalog #: 11547-015"
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organism="Homo sapiens"
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Insert Length: 502 Std Error: 0.00
Seq primer: -40UP from Gibco
High quality sequence arm.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                             cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.B. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumor Gene Index
Unpublished (1997)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/note="Organ: lymph node; Vector: pCMV-SPORT6; Site_1: Sal1; Site_2: Not1; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.25 kb. Life Technologies catalog #: 11547-015"
                   /tissue_type="serous papillary carcinoma,
pooled tumors"
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/db_xref="taxon:9606"
/clone="IMAGE:2134162"
                                                            /organism="Homo sapiens"
/mol type="mRNA"
/db xref="taxon:9606"
/clone="IMAGE:2181833"
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lab_host="DH10B"
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                                       high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Michael
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/clone_lib="NCI_CGAP_Ut4" /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sall; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT

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REFERENCE
AUTHORS
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AI538541/c
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   Alignment Scores: Pred. No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.

Email: cgapbe-r@mail.nih.gov

Tissue Procurement: Ash Alizadeh, John Byrd, M.D., Mike Grever,
M.D., Louis M. Staudt, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AI538541 19 bp mRNA linear EST 13-APR-1999 td08a11.xl NCI_CGAP_CLL1 Homo sapiens cDNA clone IMAGE:2075036 3' similar to TR:Q60843 Q60843 LUNG KRUPPEL-LIKE FACTOR; mRNA
                                                                                                                                                                                                                                                                                                                                                        High quality sequence stop: 1.
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National Cancer Institute, Cancer Genome Anat
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AI538541.1 GI:4452676
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                                                             Average insert size 1.48 kb. Life Technologies catalog 11542-016"
                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:2075036"
                                                                                                                                                                                                                                                                                                                                     ocation/Qualifiers
                                                                                                                                                                                                                          lab_host="DH10B"
                                                                                                                                                                                                                                         tissue_type="B-cell, chronic lymphotic leukemia"
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                Query Match:
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DB:
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Percent Similarity:
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 19)
NCI-CGAP http://www.nobi.nlm.nih.gov/ncicgap.
NCI-CGAP http://www.nobi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP);
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA Library Preparation: David B. Krizman, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Herbert Morse, M.D., Michael R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1997)
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                                                                                                                                                                                                               /clome_lib="NCI_CGAP_HSC3"
/note="Organ: bone marrow; Vector: pAMP1; mRNA made from lymphoid tissue, CDNA made by oligo-dT priming.
Directionally cloned. Size-selected on agarose gel, average insert size 500 bp. primary library, non-amplified. cDNA Library Preparation: David B. Krizman, Ph.D. Reference: Krizman et al. (1996) Cancer Research 56:5380-5383."
                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="IMAGE:2246456"
/tissue_type="CD34+, T negative, patient with chronic
myelogenous_leukemia"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mol_type="mRNA"
/db_xref="taxon:9606"
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  4.05e+06
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RESULT 292
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        sequence.
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                                       AI597783 19 bp mRNA linear EST 14-MAY-1999 tr92g04.x1 NCI_CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2226582 3' similar to SW:DDX3_MOUSE Q62167 DEAD BOX PROTEIN 3 ;, mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Insert Length: 3996 Std Er:
Seq primer: -40UP from Gibco
High quality sequence stop: 
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1 (Dases 1 to 19)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP);
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similar to SW:PRPL_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Life Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
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EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="addoncarcinoma"
/lab_host="DH10B"
/clone_lib="NCI_CGAP_Pan1"
/note="Organ: pancreas; Vector: pCMV-SPORT6; Site 1: Sal1;
Site_2: NotI; Cloned unidirectionally. Primer: OTigo dT.
Average insert size 1.72 kb. Life Technologies catalog #:
11548-013"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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AI664013
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       39 AlaThrSer 41
Tumor Gene Index Unpublished (1997)
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
MGI:934336
                                                                                                                                                                                                                                                                                                                                      AI664013 19 bp mRNA linear EST 10-MAY-1999 ue73all.rl Soares NMPu Mus musculus cDNA clone IMAGE:1496732 5' similar to TR:089050 089050 MUSKELIN. ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Life Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 842 Std Error: 0.00
                                                                                                                                                                           Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 19)
                                                                                                                                                                                                                                           Mus musculus
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AI664013.1 GI:4767596
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Seg primer: -40UP from Gibco
High quality sequence stop: 1
                                                                                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                              Mus musculus (house mouse)
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Unpublished (1997)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol type="mRNA"
/db wref="taxon:9606"
/clone="IMAGE:2226582"
/tissue_type="adenocarcinoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clome_lib="NCI_CGAP_Pan1"
/note="Organ: pancreas; Vector: pCMV-SDORT6; Site_1: Sal1;
Site_2: Not1; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.72 kb. Life Technologies catalog #:
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AI664013/c
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                                                                                                                                                                                                                                                             MGI:934336
                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLML;
IMAGE Consortium (info@image.llnl.gov) for further
                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                         Tumor Gene Index Unpublished (1997)
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similar to
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                    sex="female"
                                            clone="IMAGE:1496732"
                                                                                                         organism="Mus musculus"
                                                                                                                                                 ocation/Qualifiers
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/note="Organ: uterus; Vector: pT7T3D-Pac (Pharmacia) with
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lab_host="DH10B"
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'sex="female"
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'db_xref="taxon:10090"
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Soares_NMPu Mus musculus cDNA clone IMAGE:1496732 5'
TR:089050 089050 MUSKELIN. ;, mRNA sequence.
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/clone lib="Soares NMPu"
/note="Organ: uterus; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from pregnant mouse uterus, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "

Ocres:
4.05e+06 Length: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1999)
Contact: Marra M/WashU-NCI Mouse EST Project 1999
Washington University School of Medicine
Washington Park Parkway, Box 8501, St. Louis, MO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Marra,M., Hiller,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T.,Gibbons,M., Pape,D., Harvey,N., Schurk,R. Ritter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and Wilson,R.
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Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MGI:995933
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AI747751.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (bases 1 to 19)
/dev_stage="embryo, 14 dpc"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                            mol_type="mRNA"
|strain="C57BL"
                                                                                                                                                                                                                                                                                                                            db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.00%
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                                                                                                                                                                                                                                                                                one="IMAGE:2088249"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    wrong strand
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US-09-966-880A-8 (1-198) x AI747751
                                                                                                                                                                                                                                                                                                                                                              Seq primer: -40UP from Gibco
High quality sequence stop: 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
Email: Ggapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.

1 (Dases 1 to 19)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AI758301.1 GI:5152024
                                                                                                                                                                                                                                                                                                                                                                                                               Trace considered overall poor quality Insert Length: 1803 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         National Cancer Institute, Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
cound through the I.M.A.G.E. Consortium/LLNL at:
                             /clome_lib="NCI_CGAP_Ut3"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site 1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.45 kb. Life Technologies catalog #:
11541-018"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     insert. Size selection was performed to exclude fragments <1.5kb</pre>
Library constructed by Dr. Sumio Sugano
(University of Tokyo Institute of Medical Science).
Custom primers for sequencing: 5' end primer
CTTCTGCTCTAAAAGCTGCG and 3' end primer
CGACCTGCAGCTCGAGCACA."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          [TGTTGGCCTACTGG], digested and cloned into distinct sites of the pME18S-FL3 vector (5' site CACTGTGTG, CACCATGTG). XhoI should be used to isolate the cDNA
                                                                                                                                                                       adenocarcinoma,
                                                                                                                                                                  /tissue_type="poorly-differentiated endometrial
/denocarcinoma, 2 pooled tumors"
                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:2278260"
                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                 lab_host="DH10B"
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Matches:
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AUTHORS
TITLE
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AI811474
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KEYWORDS
                                                 Query Match:
DB:
                                                                                           Percent Similarity:
Best Local Similarity:
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US-09-966-880A-8 (1-198) x AI811474 (1-19)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.

Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 19)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq primer: -40UP from Gibco
High quality sequence stop: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Trace considered overall poor quality Insert Length: 1729 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              element MSR1 repetitive element ;, mRNA sequence. AI811474
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                 /clone_lib="NCI_CGAP_Ut1"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.75 kb. Life Technologies catalog #:
                                                                                                                                                                                                                                                                                                                                                                               /clone="IMAGE:2262438"
/tissue type="well-differentiated endometrial
adenocarcinoma, 7 pooled tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                11538-014"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
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Matches:
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REFERENCE AUTHORS TITLE

COMMENT

JOURNAL

Tumor Gene Index Unpublished (1997)

ORIGIN

FEATURES

Insert Length: 1803

source

ACCESSION VERSION

SOURCE ORGANISM

Homo

sapiens

Homo sapiens (human)

KEYWORDS

RESULT 296 AI758301/c

DEFINITION Pocus 밁 S

1

104 LeuSerLeu 106

Query Match: DB:

Percent Similarity: Best Local Similarity:

3.00 100.00% 100.00% 4.05e+06

52%

ORIGIN

Alignment Scores:

No.:

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REFERENCE
AUTHORS
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RESULT 299
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                                                                                                                                                                   Query Match:
DB:
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                                                                                                                               US-09-966-880A-8 (1-198) x AI918188 (1-19)
                                                                                                                                                                                                                           Percent Similarity:
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                                                     16
                                                                                         30 CYSTYTVAL 32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LINL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1154 Std Error: 0.00
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Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 19)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
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Unpublished (1998)
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h quality sequence stop: 1.
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/tissue_type="anaplastic oligodendroglioma"
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                                                                                                                                                                                                                                                                     AUTHORS
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1 (bases 1 to 19)

Brenner; S., Williams; S.R., Vermass; E.H., Storck, T., Moon, K., McCollum, C., Mao, J.I., Kirchner, J.J., Eletr, S., DuBridge, R.B., Burcham, T. and Albrecht, G.

In vitro cloning of complex mixtures of DNA on microbeads: Physical separation of differentially expressed cDNAs

Broc. Natl. Acad. Sci. U.S.A. 97 (4), 1665-1670 (2000)
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Dictyostelium discoideum
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PROJECT = Dictyostelium discoideum cDNA project in Japan.
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Tel: 81-298-53-4664
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1 (bases 1 to 19)
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                                   LYNX Therapeutics, Inc.
25861 Industrial Blvd., Hayward, CA 94545, USA
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510 670 9302
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Sequence obtained from LYNX Therapeutics Megasort technology.
Collected from the up-regulated gate.
High quality sequence stop: 19.
Location/Qualifiers
1. .19
                                                            3 ACTTTGGGA 11
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Aas29280	Aaf80665	Aaa46974	Aac60546	Aaa38137	Aax87306	Aaz37511	Aav68469	Aaq71065	Acf03639	Aah58208	Aah58207	Aai65672	Aaa83046	Aaa83045	Aat41855	Aaz44774	Abz61384	Abz61946	Ada99328	Ada99324	Ada99327
Human mdm	Human mdm	Probe use	Human fra	Polynucle	PROS09 re	Human mdm	Oligo con		Huma	Cel:	Cell-cycl		cdk6 ribo		Probe/pri	Human	Human H-R	Human H-R	Human MDZ	Human MDZ	Human MDZ

ALIGNMENTS

RESULT 1 AAA20708

AAA20708 standard; RNA; 17 BP.

AAA20708;

19-JUN-2000 (first entry)

Integrin alpha 6 subunit substrate sequence SEQ ID NO:3934

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss.

Homo sapiens.

WO9950403-A2.

07-OCT-1999.

24-MAR-1999; 99WO-US006507

27-MAR-1998; 98US-0079678P.

(RIBO-) RIBOZYME PHARM INC.

Pavco PA, Roberts E, Jarvis T, Coeshott C, Mcswiggen JA;

WPI; 1999-591315/50

Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors.

Claim 55; Page 162; 305pp; English.

The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNIT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA177167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT,

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RESULT 2
AAZ75674/c
ID AAZ75674/c
AAZ75674/c
AC AAZ756
AC AAZ656
CC Invent
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23-NOV-1998;
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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification

Novel :

Page 2369; 2745pp;

English.

This sequence represents an antisense oligonucleotide that targets genes of the phosphoenolpyruvate-sugar phosphotransferase system (PTS) operon. The invention relates to antisense oligonucleotides (e.g. the present sequence) which interact with and inhibit translation of a target RNA sequence in a bacterium. The RNA sequences that the oligonucleotides target encode proteins such as enzymes for biosynthesis of cell wall proteins, ribosomal RNA, ribosomal proteins, proteins essential for nutrient uptake, proteins associated with pathogenicity, subunits of DNA dependent RNA polymerase, and DNA polymerase. The antisense

Claim 10;

Page 29; 50pp;

English.

biallelic markers used to construct a high density disequilibrium

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16-SEP-1998;
16-SEP-1998;
16-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primers for the biallelic markers. The biallelic markers of the invent have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studie which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the sequence in the sequence is the sequence of the marketing from the sequence of the sequence of the sequence of the sequence is the sequence of the sequence in the sequence is sequence.
                                                                                                                                                                       New antisense oligonucleotide, useful for treating and diagnosing bacterial infections, interacts with and inhibits translation of RNA sequence in bacteria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide; treat; inhibit translation; diagnose; cell wall biosynthesis; ribosomal RNA; ribosomal protein; pathogenicity; nutrient uptake; bacterial infection; PTS operon; Haemophilus influenzae;
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CC clearing activity, which specifically cleave RNA encoded by an aryl complete the control of t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>..</u>
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នននិន្ននិង
Best Loc
Matches
                                 Query Match
                                                                                                     macular degeneration (ARMD), inflammation, and arthritis, as well neovascular glaucoma, myopic degeneration, psoriasis, verruca vul angiofibroma of tuberous sclerosis, por wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu synd and other syndromes and diseases related to the levels of ARNT, Integrin subunit alpha-6, or integrin subunit beta-3
                                                                     Sequence 17
               Local
                  Similarity
                                                                     BP; 8 A; 3 C;
                2.3%;
78.6%;
                                                                       N
                                                                     G; 0 T; 4 U; 0 Other;
               Score 14; DB 2;
Pred. No. 1.2e+04;
                               Length 17
                                                                                                                                                                                               as well
                                                                                                                                             syndrome,
                                                                                                                                                                               vuigaris,
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37 TACCAATTCAAAAA
Ь
UACCAAUUCAAAAA
                50
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Conservative

₩ --

Mismatches

0

Gaps

0

밁 Ś

12-JUN-2003 ABT34570; ABT34570 standard; DNA; 17 suppression (first entry) related human fukutin oligo SEQ Ħ ŏ

schizophrenia; human fukutin; Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; protein chip; gene ds. degeneration; therapy; tumour suppression;

WO2003025175-A2 Homo sapiens

27-MAR-2003

17-SEP-2002; 2002WO-IB004208.

17-SEP-2001; 2001FR-00011978

(MOLE-) MOLECULAR ENGINES LAB

Telerman A,

Amson

æ

Tuijnder M;

2003-313353/30

New isolated nucleic acid, useful for treating viral diseases with tumors and cell degeneration, also related polypeptides, and transfected cells. associated antibodies

Disclosure; Page 58; 720pp; French

RESULT 5
ABT34570
ID ABT34570
ID ABT3
XX ABT3
XX ABT3
XX ABT3
XX ABT3
XX Cyto
XX Cyto
XX Cyto
XX Schill
KW Schil
KW Schil
KW Schil
KW Schil
KW Homo
XX The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the rector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell degeneration. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and

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RESULT 6
ADB42497
ID ADB42497
XX ADB4

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                      The invention relates to the isolation of 6327 nucleotide sequences, all fragments of at least 180% identify, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding NNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro sense and antisense sequences, of nucleotides involved in tumour suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as experimental models. The nucleotides (also vectors containing them and cells containing the vectors), the encoded polypeptides and antibodies ((Ab) against the polypeptide are useful for prevention and/or treatment of viral infections or diseases characterized by development of tumours or cell degeneration (e.g. Alaheimer's disease or schizophrenia).

Chalysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can be used to screen for their specific interactive molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polymucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-DEC-2003
04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer; probe; tumour suppression; tumour reversion; apoptosis;
virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB42497 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 361; 771pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-SEP-2001; 2001FR-00011981
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2003-441574/41
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Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antiviral; neuroprotective; nootropic; neuroleptic; ss;
   d to screen for their useful for treating of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
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GGGCTGCGGCGGCT 386

Query Match Best Local : Matches 1

Similarity

2.3%;

Score 14; Pred. No.

DB 2; L

Length 20

Conservative

0

Mismatches

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SXS
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Matches
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                                   This is a novel oligomer comprising 20 covalently linked nucleotides which bind to the 5' Cap region of the interleukin-1 receptor (ILIR) mRNA. Expression of ILIR, in cells and tissues can be modulated by compositions comprising oligomers which are able to specifically hybridise with target areas of its encoding sequence. The composition can be used for treatment of disease in humans caused by excessive receptor expression, e.g. inflammation. When labelled they can be used diagnostically to determine overexpression of ILIR, also to determine localisation and distribution of this expression for research, diagnostic
                                                                                                                                                                                                     2'-substituted oligonucleotide(s) specific for interleukin-1
type I - used to modulate expression and detect overexpression
                                                                                                                                                                                                                                                                                                                      21-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                        27-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Type I interleukin-1 inflammation; ss; 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Phosphorothiate oligonucleotide sequence 8051 targeting ILIR mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression of the nucleotides
  Sequence
                                                                                                                                                                                                                                           WPI; 1998-018646/02.
                                                                                                                                                                                                                                                                    Miraglia L,
                                                                                                                                                                                                                                                                                                                                             12-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                 WO9744656-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                          or therapeutic purposes
                                                                                                                                                               Example 5; Page 19; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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14; Conserv
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ilarity 100.0%;
Conservative
 BP; 1
                                                                                                                                                                                                                                                                    Bennett CF,
                                                                                                                                                                                                                                                                                              PHARM INC
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                                                                                                                                                                                                                                                                                                                      96US-00651692
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/note= "Phosphorothioate internucleotide linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
 A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptor; IL1R; human; IL1 protein; hybridisation; Cap region; phosphorothicate linkage.
 8 C; 8
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 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                    z
                                                                                                                                                                                                                                                                   Geiger T;
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1.2e+04;
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GGGCTGCGGCGGCT 20

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RESULT 9
ABZ30588/c
ID ABZ305
XX
AC ABZ305
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DT 30-JAN
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                                                                                                                                                                                                                                          This is a novel oligomer comprising 20 covalently linked nucleotides which bind to the 5' Cap region of the interleukin-1 receptor (ILIR) mRNA. Expression of ILIR, in cells and tissues can be modulated by compositions comprising oligomers which are able to specifically hybridise with target areas of its encoding sequence. The composition can be used for treatment of disease in humans caused by excessive receptor expression, e.g. inflammation. When labelled they can be used diagnostically to determine overexpression of ILIR, also to determine localisation and distribution of this expression for research, diagnostic
                         ABZ30588;
                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                           2^\prime\text{-substituted} oligonucleotide(s) specific for interleukin-1 receptor type I - used to modulate expression and detect overexpression of the
                                               ABZ30588
                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 19; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Miraglia L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-018646/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Phosphorothiate oligonucleotide sequence 8054 targeting ILIR mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Type I interleukin-1 receptor; IL1R; human; IL1 protein; hybridisation; inflammation; ss; 5' Cap region; phosphorothicate linkage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-JUN-1998
                                                                                                                                                                         Local
                                                                                                                                                                                                                                    therapeutic purposes
                                                                                                                                     373
                                                                                                                                                             14;
                                                                                                             N
                                                                                                                                                                         Similarity
                                                standard;
                                                                                                           GGGCTGCGGCGGCT 15
                                                                                                                                    GGGCTGCGGCGGCT 386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
                                                                                                                                                             Conservative
                                                                                                                                                                                                          BP; 2 A; 7 C; B G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bennett CF,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                DNA;
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                                                                                                                                                                        Score 14;
; Pred. No.
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                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Geiger
                                                                                                                                                                                 DB 2; Length 20;
                                                                                                                                                                        1.2e+04;
                                                                                                                                                            Indels
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                                                                                                                                                           Gaps
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30-JAN-2003

(first entry)

RESULT 10
ACA90226
ID ACA90
XX
AC ACA90
XX

ACA90226 standard;

DNA; 20

ACA90226;

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Query Match Best Local Similarity

2.3%;

Score 14; Pred. No.

DB 6; Length 20, 1.2e+04;

Matches

Conservative

0;

Mismatches

Indels

0

Gaps

0

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cc is essential to the survival or growth of a fungus, a gene that contributes to the virulence and/or pathogenicity of a fungus, a gene ct that contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a diploid fungus and for identifying a therapeutic agent for treatment of a mammalian considerable of a gene product, preferably enzymatic activity. Carbon compound catabolism, biosynthetic, transporter, transcriptional, conspound catabolism, biosynthetic, transporter, transcriptional, conspound catabolism, biosynthetic, transporter, transcriptional, constituty. The method is useful for identifying a compound having the activity. The method is useful for identifying a compound having the activity. The method is useful for identifying a compound having the ability to inhibit growth or proliferation of C. albicans cells and for treating infection by C. albicans. The present sequence is that of a PCR conserved used in the method of the invention. Note: The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                               The invention relates to constructing (MI) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying one allele by insertion or replacement by a cassette having an expressible selectable marker and modifying other allele by recombination, of a promoter replacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter. (MI) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are modified. The diploid fungal cells having both alleles modified are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that
  Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 36; SEQ ID NO 4739; 167pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-566694/60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-DEC-2000; 2000US-0259128P
20-FEB-2001; 2001US-00792024
22-AUG-2001; 2001US-0314050P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      proliferation; Candida albicans; fungicide; antifungal; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-DEC-2001; 2001WO-US049486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Candida albicans GRACE strain PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jiang B,
BP; 3 A; 5 C; 6 G; 6 T; 0 U; 0
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Other;
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RESULT ADC2651.
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09-OCT-2001; 2001US-0328056P.
12-OCT-2001; 2001US-032844P.
15-OCT-2001; 2001US-0339414P.
17-OCT-2001; 2001US-0339142P.
22-OCT-2001; 2001US-0341058P.
24-OCT-2001; 2001US-0343629P.
29-OCT-2001; 2001US-0349575P.
01-NOV-2001; 2001US-034957P.
25-UN-2002; 2002US-0391342P.
01-OCT-2002; 2002US-0391342P.
                                                                                                                                                                                                                                                                          The invention describes an isolated polypeptide comprising any of 33 90 1273 amino acid sequences (I) given in the specification or its mature form, a sequence that is at least 95 % identical to (I), or a sequence comprising one or more conservative substitutions in the amino acid sequence of (I). The polypeptide is useful for preparing a composition for treating or preventing e.g. cancer. This sequence represents a primused to isolate DNA encoding a novel human NOV protein
 NOV; cytostatic; metabolic
                          NOV protein-related
                                                18-DEC-2003
                                                                                              ADC26512 standard; DNA;
                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                           Example 20C; Page 212; 253pp;
                                                                                                                                                                                                                                                                                                                                                                                                preventing e.g., cancer.
                                                                                                                                                                                                                                                                                                                                                                                                            New DAFK3 polypeptide, useful for preparing
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-381704/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alsobrook JP, Burgess CE, Catterton E, Chant JS, Edinger SR, Gerlach VL, Giot L, Gorman L, Guo X, Mezes PS, Millet I, Ooi CE, Patturajan M, Rieger Taupier RJ, Zerhusen BD, Zhong H, Zhong M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel
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                        forward
                                               entry)
                                                                                                                                                                                                                  2.3%;
                                                                                                                                                                                                                                                     4 C; 6
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                                                                                              BP
 disorder;
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                        PCR primer SEQ ID
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                                                                                                                                                                                                                  Score 14;
Pred. No.
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immune; neurodegenerative;
                                                                                                                                                                                                                  DB 7; Lo
                                                                                                                                                                                                                                                                                                                                                                                                              a composition for treating or
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Rieger DK,
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The invention relates to a novel isolated NOV polypeptide. The polypeptide of the invention demonstrates cytostatic activity and may be used for determining the presence of, or predisposition to a disease associated with altered levels of expression of the polypeptide, including metabolic disorders, immune disorders, neurodegenerative disorders, circulatory diseases, haemopoletic disorders, wasting diseases and cancer. The polypeptide may also be utilised during gene therapy procedures, vaccine development and transgenic animal production. The current sequence is that of the PCR primer of the invention which was
                                                                                                                                                                                        New isolated NOVX polypeptide, useful for determining the presence of, or predisposition to a disease associated with altered levels of expression of the polypeptide, and for treating or preventing cancer.
                                                                                                                                                                                                                                              WPI; 2003-221607/21.
                                                                                                                                                                                                                                                                                  Padigaru M,
Shimkets RA,
                                                                                                                                                                  Example C;
                                                                                                                                                                                                                                                                                                              Guox,
                                                                                                                                                                                                                                                                                                                          Catterton
                                                                                                                                                                                                                                                                                                                                     Anderson DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-MAR-2002;
02-APR-2002;
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01-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          circulatory; haemopoietic; wasting; cancer; gene therapy; vaccine;
transgenic; human; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                 16-APR-2002;
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                                                                                                                                                                                                                                                                   son DW, Berghs C, Boldog FL, Bur
irton E, Edinger S, Eisen AJ, Ell
; Jeffers M, Kekuda R, Li L, Ma
jaru M, Patturajan M, Pena CEA, R
ets RA, Spaderna SK, Spytek KA,
it CAM, Voss EZ, Zhong M;
                                                                                                                                                                                                                                                                                                                                                                 CURAGEN CORP.
                                                                                                                                                                SEQ ID NO 337; 478pp; English.
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2002US-0361677P.
2002US-0363637P.
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2002US-0360814P.
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2002US-0345038P.
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2001US-0308877P
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Ellerman K, Ger
, Malyankar UM,
A, Rastelli L, S
KA, Stone DJ, Ta
                                                                                                                                                                                                                                                                                 Casman SJ;
Gerlach V, Go
M, Miller CE;
Shenoy S;
Taupier RJ;
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Query Match Best Local Similarity

2.3%; 100.0%;

Score 14; Pred. No.

DB 9; L

Length

20

Sequence

20

BP; 2 A; 6 C; 5 G; 7 T; 0 U; 0 Other;

human NOV DNA.

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RESULT 13
ABK24744
ID ABK24
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ABT05327
ID ABT05
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Best Local
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                                                                                                                                                      The invention comprises the amino acid and coding sequence of the human N -acety/galactosaminidase (NAGA) alpha protein. The invention specifically comprises novel polymorphic sites identified within the NAGA gene. The NAGA gene is located on chromosome 22q13.2-q13.31, and encodes a lysosomal glycohydrolase that cleaves alpha-N-acety/galactosaminyl moieties in glycoconjugates. The NAGA mad protein sequences of the invention are useful for studying the expression and function of NAGA and for screening candidate drugs to treat diseases related to NAGA activity. The NAGA gene polymorphisms identified in the present invention are useful for haplotyping and genotyping the NAGA gene of an individual. The present DNA sequence represents an N-acety/galactosaminidase gene allele-
ABK24744
                ABK24744 standard; DNA; 17
                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                         WPI; 2002-566449/60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; PCR; primer; ss; gene therapy; N-acetylgalactosaminidase alpha; chromosome 22q13.2-q13.31; lysosomal glycohydrolase; screening; SNP; NAGA-related disease; single nuclectide polymorphism; haplotyping; NAGA;
                                                                                                                                                                                                                                                                     Claim 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABT05327 standard; DNA; 15
                                                                                                                                                   specific
                                                                                                                                                                                                                                                                                      function
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human N-acetylgalactosaminidase
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                                                                                                                                                                                                                                                                                                                                                             (GENA-)
                                                                                                       Local
                                                                                                                                                                                                                                                                                   genetic variants of isolated N-acetylgalactosaminidase (NAGA), e, useful for therapeutic purposes, for studying the expression ction of the polymucleotide, and for expressing NAGA protein.
                                                                              174
                                                                                                                                                                                                                                                                                                                                                                                                                                                    sapiens.
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                                                                             ATTGCTCTTCCTC 186
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                                                                                                                                                                                                                                                                                                                                         Kazemi A,
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llarity 100.0%;
Conservative (
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                                                                                               Mismatches
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                                                                                                                                 T; 0
                                                                                                                                U; 1 Other
                                                                                                      DB 6; Le
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                                                                                                               Length 15;
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Best Local Similarity
Matches 13; Conserv
                                        Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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    conservative 0;
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CC respect to the genetic sequence to be altered and further comprises
CC chemical modifications of the oligonucleotide. The chemical modifications
CC consist of o-methyl modification, an LNA modification, two or more
CC phosphorothicate linkages on a terminus, or a combination of any two or
CC directing repair or alteration of plant genetic information. The
CC oligonucleotides are particularly useful for creating plants with desired
CC phenotypes, e.g. environmental or abiotic stress tolerance, improved
CC phenotypes, e.g. altering amino acid content of plants or
CC conferring amino acid over production), herbicide resistance (e.g.
CC glyphosste resistance, imidazolinone and sulphonylurea herbicide
CC resistance, porphyric herbicide resistance or triazine resistance),
CC disease resistance, modified oil production, modified starch production
CC (e.g. increased starch or production of waxy starch), altered floral
CC (e.g. reduced palmitate, increased stearch or reduced linolenic acid).
CC The oilgonucleotides are also useful for producing albino mutants for the
CC analysis of photosynthetic processes. This sequence represents a genome
CC altering oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA repair; DNA alteration; environmental tolerance; hygromycin-B; abiotic stress tolerance; improved nutritional value; hygromycin; primer amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; triazine resistance; disease resistance; modified oil production; modified strarch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced palmitate production; albino plant; increased stearate production; reduced linolenic acid production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an oligonucleotide for targeted alteration genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New oligonucleotides with modified nuclease-resistant termini, usef creating plants with desired phenotypes, e.g. stress tolerance, imp nutritional value, herbicide or disease resistance, or modified oil
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30-OCT-2000; 2000US-0244989P
27-MAR-2001; 2001US-00818875.
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phosphorothicate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphonylurea herbicide resistance, porphyric herbicide resistance or triazine resistance), disease resistance, modified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced linolenic acid).
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30-OCT-2000; 2000US-0244989P.
27-MAR-2001; 2001US-00818875
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                                                                                                                                                                                                                                                                                                                                                                                                         genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with having a DNA domain. The DNA domain has at least one mismatch with respect to the generatic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modification consist of onethyl modification, an LNA modification, two or more the property of the consist of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New oligonucleotides with modified nuclease-resistant termini, useful creating plants with desired phenotypes, e.g. stress tolerance, impronutritional value, herbicide or disease resistance, or modified oil
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Best Local S
Matches 13
has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene we mapped to human chromosome 10p12.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
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09-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                                                 protein
                                                                                                                                                                                                                                                                                                                                                                                       The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 168; 718pp; English.
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                                                                                                                                                                                                                                                                                                                                                              sent invention relates to human testis expressed Patched like (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL
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2001US-0327898P.
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Pred. No.
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Matches 13
The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10pl2.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
                                                                                                                                                                                                                                                                                                                                                              Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-JAN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          foetal liver, bone marrow, brain, k skeletal muscle or colon function. clinically useful diagnostic marker male infertility and cancer. The prexample from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-MAY-2001;
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; 2001WO-US000664.
; 2001WO-US000665.
; 2001WO-US000667.
; 2001WO-US000668.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       suppressor;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 T; 0 U; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HTPL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids ar clinically useful diagnostic markers and potental therapeutic agents male infertility and cancer. The present oligonucleotide was used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-OCT-2001;
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23-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AEOM-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  muscle; colon; male infertility;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   먥
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer; ss.
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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV7875 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similate to that of patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene we mapped to human chromosome 10p12.1. HTPL and its coding sequence are Novel isolated human testis expressed Patched like protein for identifying agonist and antagonist and specific binding for treating subjects having defects in HTPL. Example 2; Page 168; 718pp; English. (HTPL), useful g partners, and similar was

2002-676582/73

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Best Local S
Matches 13
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30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-WAY-2001; 2001US-00864761.
                                       The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention
shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is
                                                                                                                                                                            Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and
                                                                                                                                                                                                                           WPI; 2002-676582/73
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                                                                                                                                                                                                                                                                                    (AEOM-)
                                                                                                                                                                                                                                                                                                                                                                                                                     30-JAN-2001; 2001WO-US000663
                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human testis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy;
                                                                                                                                    2;
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                                                                                                                                                                                                                                                                                    AEOMICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ne therapy; tumour suppressor; HTPL; chromosome 10p12.
tis expressed Patched like protein; testis; adrenal; 1
cell development; bone marrow; brain; kidney; lung; p
skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                    Page 168;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              scanning
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                                                                                                                                  English
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                                                                                                                                                                 defects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                 'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 6; Le
3.8e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chromosome 10p12.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                         TALH
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RESULT 19
ABV79550/c
ID ABV795
XX ABV795
XX ABV795
XX O3-JAN
XX Human;
KW Human;
KW Prosta
XX O7-AUG
XX Prosta
XX O7-AUG
XX Prosta
XX O7-AUG
XX O7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-676582/73.
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23-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         prostate; skeletal muscle; colon; male infertility; cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABV79550 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   male germ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                testis expressed Patched like protegerm cell development; bone marrow;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HTPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AEOMICA INC
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ed Patched like protein; testis; adrenal; liver;
opment; bone marrow; brain; kidney; lung; placen
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Pred. No.
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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The

Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.

Example 2;

Page 168; 718pp;

English

ភ្នំពីស្តីស្តីស្តីស្តីស្តីស្តីស្តី ស្ត្រីស្តី ស្ត្រីស្តី ស្ត្រីស្តី ស្ត្រីស្ត្រីស្ត្រីស្ត្រីស្ត្រីស្ត្រីស្ត្រី

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RESULT 20
ABL31517
ID ABL31
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Matches 13;
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Best Local
The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonuclectides (ABL3012-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnossis of genetic diseases and identifying individuals
                                                                                                                                                                                                                                                                                                                                                                                                                              Human leukocyte antigen (HLA) ty individuals e.g. by determining transplanting between them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10pl2.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                          Claim 10; Page 284; 345pp; Japanese.
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                                                                                                           This sequence represents an isolated nucleic acid sequence associated with tumour suppression or regression, apoptosis or virus resistance. invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to diagnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
                                                                                                                                                                                                                              New nucleic acid sequences associated with tumor suppression, rappoptosis or virus resistance are useful to diagnose and treat disease, development of tumor cells and cell degeneration.
                                                                                       Sequence 17
                                                                                                                                                                                                        Claim 1; Page 512;
                                                                                                                                                                                                                                                                                                     Tuijnder M,
                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ss; tumour suppressor; antitumour; cytostatic; tumour suppression; tumour regression; apoptosis; virus resistance; diagnosis;
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RESULT 23
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Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The mucleic acids can also make the first of the model 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of
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                     Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21,3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
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                                                                                                                                Human MDZ3 scanning oligonucleotide SEQ ID 314.
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12; chromosome 7q22.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 5g21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15g26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy,
                                                                                                                                                            Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7 chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q
                                                                                                                                                                                                                                                                         20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 3 A; 4 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shannon M, Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EP1281758-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                           Homo sapiens
                                                                                                                                              developmental
                                                                                                                                                                                                                                   Human MDZ3 scanning oligonucleotide SEQ ID 316
                                                                                                                                                                                                                                                                                                                                                ADA99327 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; SEQ ID NO 314; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          246 CTCCTGGAGCCCC 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                              disorder; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13;; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 7; Length 17; 3.8e+04;
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30-JUL-2002; 2002EP-00016874

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC The present invention relates to novel human zinc finger-containing CC proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is CC encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, CC MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome (cc 15q26.1). The MDZ3, MDZ4, MDZ7, and MDZ12 is encoded at chromosome (cc 15q26.1). The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, CC or in manufacturing a medicament for treating or preventing a disorder CC associated with decreased or increased expression or activity of MDZ3, CC MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic CC acids and proteins are also useful for diagnosing or monitoring a disease CC caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic CC acids can also be used as probes to detect and characterize gross CC alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are Useful in constructing microarrays for measuring gene expression. The CC proteins are useful as therapeutic agents for gene therapy or as CC vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Loc
Matches
WPI; 2003-423107/40
                                 Shannon M,
                                                                                                                                 30-JUL-2002; 2002EP-00016874
                                                                                                                                                                   05-FEB-2003
                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                       developmental disorder; ss.
                                                                                                                                                                                                                                                                              Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p11.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
                                                                                                                                                                                                                                                                                                                                                      Human MDZ3 scanning oligonucleotide SEQ ID 313.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17
                                                                                                02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                           20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA99324 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-AUG-2001; 2001US-00922181.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTCCTGGAGCCCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTCCTGGAGCCCC 258
                             Gu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 316; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 2 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                             Nguyen C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.2%; Score 13;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 7; Lc...
o. 3.8e+04;
o;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
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New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
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Example 8; SEQ ID NO 313; 103pp; English.

proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 1pg1.2 and MDZ12 is encoded at chromosome 1pg26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 is encoded at chromosome 1pg26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as The present invention relates to novel human zinc finger-containing The present sequence was used to illustrate the invention.

S Matches Query Match Local 246 CTCCTGGAGCCCC 258 2.2%; Similarity 100.0%; Conservative 0 Score 13; Pred. No. Mismatches DB 7; L; 3.8e+04; 0 Length 17 Indels 0, Gaps 0

Sequence 17 BP; 3 A; 3 C; 8 G; 3 T; 0 U; 0 Other;

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CTCCTGGAGCCCC 5

RESULT 26
ADA99328/c
ID ADA993
XX ADA993
XX ADA993
XX 20-NOV
XX Cytost
KW Zinc f
KW Zinc f
KW Zinc f
KW Chromo
KW develo
XX Homo s
XX EP1281
XX D5-FEB
XX O5-FEB
XX NEP1281
XX NEP1281
XX NEP1281
XX NEP1281
XX AEOMXX WPI; 2
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XX WPI; 2
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PI masoci Human MDZ3 scanning oligonucleotide SEQ ID 317. 20-NOV-2003 ADA99328 ADA99328 standard; DNA; 17 (first

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2

05-FEB-2003

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC

Shannon M, Gu Y, Nguyen C;

2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer. of MDZ3

Example 8; SEQ ID ö 317;

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RESULT 27
ABZ61946/c
ID ABZ619
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are
                                                                                                                                                                                                                                                                   Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2001; 2001US-0294140P
06-JUN-2001; 2001US-0296249P
10-SEP-2001; 2001US-0318471P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                                                                       Claim 58;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME
                                                                                                                                                                                                                                                                                                                                                                                 2003-140484/13
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                                                                                                                                                                                                                    Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 2 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                 125; 185pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM INC
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.8e+04;
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В Ś

treating

ovarian,

colorectal,

lung, prostate

Query Match
Best Local Similarity
Matches 12; Conserv

2.2%;

<u>.</u>.

Score 13; DB 7; Pred. No. 3.8e+04; l; Mismatches (

0; Indels Length 17;

0

Gaps

0

Sequence 17

BP; 0 A; 5 C; 11

G; 0 T; 1 U; 0 Other;

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RESULT 28
ABZ61384
ID ABZ61
XX ABZ61
XX ABZ61
XX ABZ61
XX Human
XX Human
XX Human
XX Homo
XX Clair
XX RIBC
XX 
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                                                                                  acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ5989 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66550 - ABZ66585 represent substrate/target sequences for the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 1 A; 6 C; 5 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 58; Page 114; 185pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200297114-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ61384;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ61384 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H-Ras DNAzyme target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15 GCAAGGCTGAGCC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCAAGGCTGAGCC 368
                                                                 of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     #175.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 7; L
3.8e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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372

GGGCTGCGGCGG

384

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RESULT 29
AAZ44774/c
ID AAZ447
XX AAZ447
XX AAZ447
XX Human
XX FADD;
KW FADD;
KW FADD;
KW POODE;
XX US6015
XX This in of the XX Usice CC nucled CC especic CC e
ARESULT 30
AAT41855/c
ID AAT418
XX AAT418
AC AAT418
XX 21-FEB
XX Probe/
XX Plasmo
KW Plasmo
KW identi
XX V
OS Synthe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel antisense oligonucleotides (OGNS) (I) 8-2 nucleotides in length that specifically hybridize with and inhibit nucleic acids encoding human Fas-associated death domain (FADD), targete to the 3' untranslated region (3'UTR). (I) can be used to treat animals, especially humans, suspected of having or being prone to a disease or condition associated with FADD expression. AAZ44746-Z44831 represent primers and probes used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotides, useful death domain (FADD) expression are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ44774 standard;
    Synthetic
                                                                                                                                                                                                      AAT41855;
                                                                                                                                                                                                                                               AAT41855 standard; cDNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 6 A; 5 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 16; Col 53-54; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monia BP,
                                      Plasmodium falciparum; erythrocyte membrane protein; malaria; detection; identification; treatment; prevention; parasite; ss.
                                                                                                           Probe/primer for Plasmodium falciparum erythrocyte membrane protein
                                                                                                                                                         21-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FADD; human; antisense; inhibitor; Fas-associated death domain; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human FADD primer ISIS #23874.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ44774;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-JAN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                        130
                                                                                                                                                                                                                                                                                                                                                             14
                                                                                                                                                                                                                                                                                                                                                                                                                                                 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                             CTGGACTTTGGTT 2
                                                                                                                                                                                                                                                                                                                                                                                                    CTGGACTTTGGTT 142
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                               ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-00357072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-00357072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.2%; Score 13;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>1</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Baker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              뭠
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for inhibiting human Fas-associated targeted to the 3' untranslated region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 3; Length 18; 3.8e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (I) 8-20
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RESULT 31
AAA83045/c
ID AAA830
XX AAA830
XX AAA830
DT 04-DEC
XX Ribozy
XX Ribozy
XX Mammal
XX WO2000
XX WO2000
XX MO2000
XX TITIZ
XX IMMU-
PF 06-DEC
XX O4-DEC
XX IMMU-
PR 1: Z
XX WPI; 2
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    reverse the sequestration of erythrocytes in patients suffering from malaria. Nucleic acids derived from the PERMP1 gene can be used as probes and primers to identify a Plasmodium falciparum parasite, the primers used to generate characteristic amplification patterns from different P. falciparum strains. Antibodies specifically immunoreactive with the PERMP1 polypeptide or its fragments may be used in diagnosis of malaria infection. Nucleic acid fragments of at least 15 contiguous nucleotides of the PERMP1 gene are also claimed. They may be generated by amplification with the probes/primers described in AAT41854-T41867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A polypeptide comprising a Plasmodium falciparum (Pf) erythrocyte membrane protein 1 (PfEMP1) or active fragments or analogues of that protein can be used in the treatment or prevention of symptoms of a malaria parasite infection. The polypeptides can inhibit, block or
                                            Tritz R,
                                                                                                                            04-DEC-1998;
                                                                                                                                                                  06-DEC-1999;
                                                                                                                                                                                                          08-JUN-2000
                                                                                                                                                                                                                                                                                          Mammalia
                                                                                                                                                                                                                                                                                                                              Ribozyme;
                                                                                                                                                                                                                                                                                                                                                                      cdk6 ribozyme binding site #105.
                                                                                                                                                                                                                                                                                                                                                                                                                 04-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA83045;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA83045 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 8 A; 2 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 24; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New Plasmodium falciparum erythrocyte membrane proteins - used to develop products for the diagnosis, treatment or prevention of malaria parasite
                                                                                                                                                                                                                                                  WC200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-497376/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-APR-1996;
                                                                                    (-DWMI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AFFY-) AFFYMAX TECHNOLOGIES NV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9633736-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15 GTTCGTCTCCA 3
                                                                                    IMMUSOL INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTTCGTCTCCA 520
                                                                                                                                                                                                                                                                                                                            hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss
                                          Welch PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pasloske
                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                          98US-0110954P
                                                                                                                                                                  99WO-US028772.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96WO-US005798
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                                            Barber JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BĽ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Howard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13; DB 2; L
Pred. No. 3.8e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                            Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               묎
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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WPI; 2000-412314/35

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AAA83046/c
ID AAA830
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1.

Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endomuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1.

Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endomuclease activity and hence is efficient in
                                                                                                                                                                                                                    New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ribozyme;
                                                                                                                                                                         Disclosure; Page 55; 109pp; English.
                                                                                                                                                                                                                                                                                                                 WPI; 2000-412314/35
                                                                                                                                                                                                                                                                                                                                                        Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                              04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hairpin; hammerhead; gene therapy; vasotropic; restenosis;
                                                                                                                                                                                                                                                                                                                                                        Welch PJ,
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                                                                                                                                                                                                                                             for inhibiting restenosis, cleaves dependent kinase other than CDK1,
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3.8e+04;
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                                                                                                                                                                                                                                               Primers AAI65647-78 were used to characterise biallelic polymorphic markers in the IBD1 gene region. The IBD1 gene encodes an inflammatory bowel disease 1 (IBD1) polypeptide, which is associated with intestinal inflammatory disease. The specification also describes a polypeptide which is in proximity to IBD1, and is designated IBD1prox. The IBD1 gene is probably involved in regulation of apoptosis and activation of NF-cc kappa B. The IBD1 and IBD1prox polymucleotides are is useful as source of probes and primers, as source of (anti)sense oligonucleotides, for recombinant production of polypeptides, and in screening for interactive compounds. The polypeptides are used to raise specific antibodies which useful for diagnostic detection or purification of IBD1 and IBD1prox, to screen for specific binding agents, potential therapeutic agents. The IBD1 and IBD1prox polymucleotides are useful for creatment and prevention of inflammatory and/or immune diseases or creatment and prevention of inflammatory and/or immune diseases or cancer, where associated with mutations in genes corresponding to IBD1 and IBD1prox, especially cryptogenetic inflammation of the intestines
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                                                                                                                                                                        Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New human nucleic acids associated with intestinal inflammatory disease, useful for diagnosis, prognosis and control of these diseases, also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAI65672;
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inflammatory disease; immune disease; cryptogenetic inflammation;
nemorrhagic rectocolitis; Crohn's disease; Blau syndrome; PCR primer;
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                                                        l Similarity
13; Conserv
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  TCCGCTGGGCTAA 65
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thes 0;
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                                                                                                                                                                                 Sequence 19
                                                                                                                                                                                                                                                                                                                                dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing such as keloid, adhesion and hypertrophic or hypertrophic burn are always and always are analyzed as the same and always and and always and always and always and always are always as the same and always and always are always as the same and always and always are always as the same always are always as the same and always and always are always as the same always and always are always and always are always as the same always are always as a same always are 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, and the control of the contr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic antipporiatic; dermatological; antiseborrheic, antidiabetic; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 117; 408pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-2001
                                                     Local Similarity
                                                                                                                                                                                                                                                        AAH57577 to AAH62099 represent sequences used in the lification of the present invention
    13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    dependent kinase cdk6 ribozyme binding site SEQ ID NO:631.
        Conservative
                                                                                                                                                                    BP; 2 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tritz R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                            100.0%;
                                                                                             2.2%;
        0
                                                Score 13;
; Pred. No.
        Mismatches
                                                                                        DB 5;
                                                3.8e+04
                                                                                    Length 19;
        Indels
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Gaps
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481

GCCTGGGAAGGGC 493

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Query Match
Best Local Similarity
Matches 13; Conserv

100.0%;

0

Mismatches

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Gaps

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2.2%; Score 13; DB 5; Louis 100.0%; Pred. No. 3.8e+04;

Length 19 Indels

Sequence

19

BP; 2

A; 7 C; 6 G; 4 T; 0 U; 0 Other;

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RESULT 35
AAH58208/c
ID AAH582
XX Cell-c
XX Feeding For Colling For Co
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                                                                                                                             cc skin or eye disease and scarring. The method involves administering a cribozyme (I) which cleaves RNA encoding a cytokine involved in the method involved in the control of the control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Treating proliferative skin or eye diseases and scarring, using ribothat cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                             scar. AAH57577 to AAH62099 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnera: proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 118; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-300427/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Robbins JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-OCT-2000; 2000WO-US029500.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisickling, ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cell-cycle dependent kinase cdk6 ribozyme binding site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bickle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCCTGGGAAGGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0161532P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               먪
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        using ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vulnerary;
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ACF03639/c
IID ACF03639/c
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ACF036
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Human
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Alzhei
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Alzhei
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Alzhei
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Alzhei
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Homo s
OS Synthe
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DR W02002
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ACF036
PH Gunthe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-NOV-2000; 2000US-0245317P.
07-NOV-2000; 2000US-0246871P.
08-NOV-2000; 2000US-0246871P.
26-JAN-2001; 2001US-0264389P.
26-JAN-2001; 2001US-0264423P.
29-JAN-2001; 2001US-0264799P.
ACF03547 to ACF03570 encode the human NOVX proteins (I) given in ABR57412 to ABR57435. (I) have cytostatic, cardiant, antiinflammatory, nootropic, immunosuppressive, antiallergic, haemostatic, anti-HIV, antidiabetic, antiarterioselerotic, anorettic, antiasthmatic, nephrotropic, virucide, antiarterioselerotic, anorettic, antiesthmatic, nephrotropic, virucide, antiparasitic, anticonvulsant, hypotensive, vasotropic, antiparkinsonian, vulnerary, angiogenic and antiangiogenic activities, and can be used in gene therapy and vaccines. The NOVX polypeptides and their antibodies can be used to determine the presence or absence of (I) in a sample. The NOVX polypeptides, polymucleotides encoding them, and antibodies against them, are useful in manufacturing a medicament for treating or preventing a syndrome associated with a NOVX-associated disorder such as hypertension, cardiomyopathy, atherosclerosis, cancer, diabetes, asthma, inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Spytek KA,
Guo X, Fe
Mishra V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiallergic; haemostatic; anti-HIV; antidiabetic; antiarteriosclerotic; anorectic; antiasthmatic; nephrotropic; antiarthritic; hepatotropic; neuroprotective; noctropic; antibacterial; virucide; antiparasitic; relaxant; anticonvulsant; hypotensive; vasotropic; antiparkinsonian; vulnerary; angiogenic; antianglogenic; gene therapy; vaccine; cancer; cardiomyopathy; atherosclerosis; hypertension; diabetes; inflammation; autoimmune disorder; allergy; blood disorder; AIDS; obesity; asthma; autoimmune disorder; allergy; blood disorder; AIDS; obesity; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gunther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Grosse WM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          muscular dystrophy; epilepsy; wasting disorder; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acquired immunodeficiency syndrome; nephropathy; cirrhosis; arthritis; Alzheimer's disease; Parkinson's disease; goitre; infection; stroke;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; NOVX; cytostatic; cardiant; antiinflammatory; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New NOVX polypeptide useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis or diabetes, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-140359/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   481
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fernandes ER,
1, Furtak K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCCTGGGAAGGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 316;
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Ellerman K,
Edinger SR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    g. cancer, cardiomyopatl
mapping, tissue typing
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                                                                                                                                                                                                                                                                                                                                                                                                                                              346pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 JR, Smithson G, Millet I, K, Alsobrook JP, Lepley DM, R, R, Gangolli EA, Gorman L, Vernet CAM, Tchernev VT, Baumgartner JC, Colman SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cardiomyopathy,
tissue typing or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pharmacogenomics.
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-FEB-1993;
04-MAR-1993;
19-AUG-1993;
22-DEC-1993;
The sequences given in AAQ71063-66 are primers which were used in a sequence-tagged site assay of the region of chromosome 22 surrounding the NF2 deletions. NF2 is a neurofibromatosis which is characterised by bilateral schwannomas. The NF2 "gene" has been shown by linkage studies to be assigned to chromosome 22. The missing or mutated gene in NF2 patients has been shown to be the merlin gene. The gene encodes a protein, merlin (moesin-erzin-radixin-like protein), which possesses tumour suppressor activity, and whose tumour suppressor activity is mediated by interactions with the cytoskeleton. The merlin gene is found on chromosome 22 between the known markers D22S1 and D22S28. The merlin gene may be used in gene therapy for the treatment of a merlin-associated tumour or NF2, or for prevention of schwannoma, meningioma, posterior
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymerase chain reaction; PCR; amplify; primer; bi-lateral schwannoma; sequence-tagged site assay; chromosome 22; NR2; deletion; hearing loss; neurofibromatosis; merlin; mossin-erzin-radixin-like protein; D22S28; tumour suppressor; activity; meningioma; cytoskeleton; gene therapy; merlin-associated tumour; D22S1; posterior capsular lens opacity; deafness; balance disorder; paralysis; ss.
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19-APR-1995
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                                                                                                                                                                                                                                                               Disclosure;
                                                                                                                                                                                                                                                                                                    The tumour suppressor gene merlin -tumours and neurofibromatosis (NF2)
                                                                                                                                                                                                                                                                                                                                                               WPI; 1994-272992/34.
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(first entry)
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93US-00108808.
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Pred. No.
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3.8e+04;
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RESULT 38
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This represents an antisense oligonucleotide to the RNA component of human telomerase (hTR) comprised in the. The invention relates to an activator-antisense complex that comprises: (a) an antisense oligo, complementary to a 12-25 nucleotide portion of the RNA component of hTR, with a hydroxyl moiety at the first end; and (b) a linker attached to the first end, and (c) an activator of RNase L attached to the linker. The activator-antisense complex may be used for inhibiting the growth of a telomerase-expressing malignant cell or tumour. The complex is used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                    New RNase L activator-telomerase antisense complex - useful
                                                                                                                                                                                WPI; 1998-609972/51.
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                                                                                                                 Example; Page 45;
                                                                                                                                           telomerase
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03-FEB-1998;
                                                                                                                                                                                                                                                                                                            13-APR-1998;
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                                                                                                                                                                                                                                  (CLEV-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   carcinoma;
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                                                                                                                                                                                                       표,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         lomerase, hTR, activator-antisense complex, malignant; enzyme; rain; tumour malignant glioma; breast tumour; renal cell cancer; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma; lymphoma; Maldenstrom's macroglobulinemia; heavy chain disease;
                                                                                                                                         activity in telomerase-expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 5 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
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98US-00018125.
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RESULT 39
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                                                                                                                                                                                                                                                                                                                                                                              hyperproliferative conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fibrosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cowsert LM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
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AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides. AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of

Example 9; Page 47; 157pp; English.

hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or

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RESULT 40
AAX87306
ID AAX87
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                                                                           Matches
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                                                                                                             Query Match
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29-APR-1998;
22-MAY-1998;
10-JUN-1998;
10-NOV-1998;
20-NOV-1998;
                                                                                                                                                                                                  This is the nucleotide sequence of reverse primer 50148.tm.r1 that can bused in the PCR amplification of DNA50148 (see AAX87265) nucleic acids coding for PRO509 (see AAY06488). This gene is amplified in various tumour lines. The invention identifies 14 genes (see AAX87254-67) that are amplified in the genome of certain human lung, colon and/or breast cancers and/or cell lines. This gene amplification is expected to be associated with overexpression of the gene product and to contribute to tumorigenesis. The encoded proteins (see AAY06477-90) may be useful targets for the diagnosis and/or treatment of of certain cancers, and materials the second of the gene and the contribute to the contr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Botstein Wood WI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    psoriasis,
                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-430385/36
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                                     N
                                                                        L Similarity
13; Conserv
                                                                                                                                                                                     predictors of the prognosis of tumour treatment
                                                                                                                                                                                                                                                                                                                                                                                             2; Page 55; 162pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  against proteins expressed in
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                                     TGGACAGCCTCTT 14
TGGACAGCCTCTT
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llarity 100.0%;
Conservative
                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                    treatment.
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98US-0083500P.
98US-0086414P.
98US-0086742P.
98US-0107783P.
98US-0109304P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tumour; diagnosis; therapy;
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                                                                                       2.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
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18
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                                                                      Score 13; DB; Pred. No. 3.8 0; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      neoplastic cells, useful
                                                                                       DB 2; Lo
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3.8e+04;
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                                                                                                           Length 20
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                                                                      Gaps
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RESULT 42
AAC60546/c
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AAA38137/c
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                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                  Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                                                             Sequences AAA38133-A38148 represent polynucleotides that hybridise with a nucleic acid sequence encoding a D-alanine:D-alanine ligase of a given species belonging to the Streptococci or Enteroccci species. The polynucleotides are used to detect bacteria belonging to the Streptococci and Enterococci genus in a sample. The polynucleotides are also used as probes or primers that are specific for particular species or groups of
                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                            species belonging to Streptococci or Enterococci genus. The oligonucleotide probes are also useful as capture probes immobilized on a substrate to capture a target nucleic acid and can be used in a detection device comprising a matrix library of probes immobilized on a substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polynucleotides derived from unknown sequences internal to the ddl genes coding for D-Alanine:D-Alanine ligase of various bacterial strains belonging to Enterococci or Streptococci genus, useful as probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Garnier F,
Casadewall
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                                                           Human fra-1 mRNA antisense oligonucleotide ISIS 109037.
                                                                                                                                       AAC60546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2; Col 57; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Streptococcus
                                   Human;
                                                                                     31-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-338486/29
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                      antiinflammatory;
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                                                                                                                                                                                                                                                     13;
                                   fra-1; antisense oligonucleotide;
                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                     GCCT
                                                                                                                                       standard;
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3, Galimand M, Courvalin P;
                                                                                                                                                                                                                                                                                                      BP; 8 A; 3 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                     Conservative
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                                                                                    (first entry)
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                      2'-methoxyethyl wing;
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                                                                                                                                                                                                                                                                 2.2%;
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                                                                                                                                                                                                                                                                 Score 13;
Pred. No.
                                                                                                                                                                                                                                                     Mismatches
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                      phosphorothicate;
2'-MOE wing; infect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Charles
                     wing; infection; cancer;
                                                                                                                                                                                                                                                                            Length
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Best Local S
Matches 13
  22-DEC-1998;
08-MAR-1999;
02-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotides which are targeted to nucleic acids encoding fra-1. The sequences may be oligodeoxyribonucleotides or chimeric oligonucleotides containing a central gap region consisting of ten 2'-deoxynucleotides, which is flanked on both sides by 2'-methoxyethyl (2'-MOE) wings. The oligonucleotides have a phosphorothioate backbone and the cytidine residues in the 2'-MOE wings are 5-methylcytidines. The fra-1 antisense oligonucleotides are useful for inhibiting the expression of fra-1 in human cells or tissues. They can be used for diagnostics, therapeutics, prophylaxis and as research reagents and in kits. Use of the antisense compounds may also be useful prophylactically, e.g. to prevent or delay inferior.
                                                                                                                                                                                                                                                                                                                                                          PRO201; PRO292; PRO327; PRO1265; PRO344; PRO343; PRO347; PRO357; PRO715; PRO1017; PRO1112; PRO509; PRO853; PRO882; tumour cell; probe; tumourigenesis; cancer; neoplastic cell growth; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel antisense compound 8-30 nucleobases in length targeted to human -1 and which specifically hybridizes with and inhibits the expression human fra-1, useful for modulating the expression of fra-1 in cells.
                                                                                                                                                                      29-JUN-2000
                                                                                                                                                                                                                        WO200037640-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Probe used to
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Synthetic.
                                                                                                                16-DEC-1999;
                                                                                                                                                                                                                                                                                                                                PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Col 41; 38pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GAACCGGAGGAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence is one of a large number of antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
  98US-0113296P.
99WO-US005028.
99WO-US012252.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            isolate cDNA encoding novel polypeptide PRO509
                                                                                                             99WO-US030095
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.2%; Score 13; DB 3; Le
100.0%; Pred. No. 3.8e+04;
... Mismatches 0;
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Best Local
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15-SEP-1999;
30-NOV-1999;
01-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoding a novel human polypeptide. The specification describes novel polypeptides designated PRC201, PRC292, PRC277, PRC11265, PRC344, PRC343, PRC3437, PRC357, PRC715, PRC61017, PRCC1112, PRC509, PRC653 and PRC682. These genes are amplified in the genome of tumour cells. The polypeptides are believed to contribute to tumourigenesis. The polypeptides are useful target for the identification of certain cancers, and may act as predictors of the prognosis of tumour treatment. Antibodies against these polypeptides are useful in the treatment and diagnosis of neoplastic cell
                                                     WPI; 2001-190948/19
                                                                        Miraglia LJ,
                                                                                                                                                                                                                                            Human mdm2
                                                                                                                                                                                                                                                                02-MAY-2001
                                                                                                                26-MAR-1998;
                                                                                                                                                           06-FEB-2001
                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                     AAF80665
                                                                                                                                                                                                                                                                                                          AAF80665 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                      26-MAR-1999;
                                                                                                                                                                               US6184212-B1.
                                                                                                                                                                                                                      Antisense; mdm2; hyperproliferation; cancer; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New anti-polypeptide antibody useful in the neoplastic cell growth and proliferation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-452188/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Botstein D,
Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primers AAA46972-73 and probe AAA46974 were used to isolate
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13; Conserv
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                                                                                                                                                                                                                                            phosphorothioate oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                     llarity 100.0%; I
Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 111; 220pp;
                                                                                              PHARM INC
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                                                                                                                                                                                                                                                               (first entry)
                                                                        Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US020111.
99WO-US021090.
99WO-US028313.
99WO-US028409.
99WO-US028301.
                                                                                                                 98US-00048810
                                                                                                                                      99US-00280805
                                                                                                                                                                                                                                                                                                                                                                                                                                              3 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                         2.2%;
                                                                         Graham
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                                                                                                                                                                                                                                                                                                          ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                   Score 13; DB; Pred. No. 3.8
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                                                                        Monia
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                                                                                                                                                                                                                                                                                                                                                                                                             DB 3; Le
3.8e+04;
                                                                        BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treatment
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Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.

Example

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RESULT 45
AAS29280/c
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-300 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro. The hyperproliferative disorder includes cancer or psoriasis
                                                             An antisense compound, useful nucleobases targeted a region of a nucleic acid encoding hun
                                                                                                                                                                                                                                                                                                                                                                                                                   Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic; anti arteriosclerotic; vasotropic; antisense; phosphorothioa
                                                                                                                                                                                                               26-MAR-1998;
26-MAR-1999;
                                                                                                                            Miraglia
                                                                                                                                                                                                                                              02-JAN-2001; 2001US-00752983
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                                                                                                                                                                                                                                                                                                                                                                modified_base
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                                                                                                                                                                                           (MIRA/)
                                                                                                                                                                      GRAH/)
                                                                                                                                                                                 (NERO/)
                                                                                                      2001-535565/59.
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                                                                                                                                                                                       MIRAGLIA L J.
                                                                                                                                                MONIA B P.
COWSERT L M.
                                                                                                                                                                   GRAHAM M J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9;
                                                                                                                                                                                 NERO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAGCCCGAGGGGC 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotide 31716.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                            Nero P,
                                                                                                                                                                                                              98US-00048810
99US-00280805
                                                                                                                                                                                                                                                                                                            methoxyethyl bases, and
                                                                                                                                                                                                                                                                                                                                            /mod
                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                   'note= "OTHER= All
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            77pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                                                                                                                                                                                                                                                                                                            _base= OTHER
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                                                                                                                            Graham
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                                                  יים (e.g. translation human mdm2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
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Pred. No.
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                                                                                                                                                                                                                                                                                                         phosphorothioate linkages,
1-6 and bases 15-20 are 2'-0-
, and bases 7-14 are deoxynucleotides"
                                                                                                                            Monia
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3.8e+04;
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                                                                      cancer, comprises
termination codon
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                                                                      codon region)
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The present invention relates to antisense compounds, 8-30 nucleobases length targeted to the 5' untranslated region, translation termination codon region, 3' untranslated region, coding region or translation star

start ä

The invention comprises a method for detecting a modification in genomic DNA, causing inactivation or alteration of a Kruppel-like factor 6 (KLF6) tumour suppressor gene. The method of the invention is useful for diagnosing, prognosing or determining the relative risk of a cancer (i.e. prostate cancer, neuroblastoma, glioblastoma, melanoma, breast cancer, ovarian cancer, head and neck squamous cell carcinoma, hepatocellular

(e.g. neuroba

DNA.

Example 7;

Page 63;

103pp; English.

15;

English

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RESULT 46
RAL42131
ID 4AL422
XX AAL422
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Best Local 9
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                                                                                                                                                                                                                                                                                                    Detecting inactivation or alteration of a Kuppel-like factor 6 (KLF6) gene, useful for diagnosing or determining the relative risk of a cancer (e.g. neuroblastoma or breast cancer) by detecting modifications of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; PCR; primer; ss; Kruppel-like factor 6; KLF6; 2AF3; tumour suppressor gene; cancer risk; prostate cancer; neuroblastoma; glioblastoma; melanoma; breast cancer; ovarian cancer; squamous cell carcinoma; hepatocellular cancer; lung cancer; colon cancer; benign hyperplasia; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful for detecting the role of mdm2 expression in various cell functions and physiological processes and useful in both clinical research and diagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense oligonucleotides of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-241784/29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             364 GAGCCCGAGGGGC 376
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Narla G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>ہ</u>
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Martignetti J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Heath
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RESULT 47
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Best Local S
Matches 13
therapeutically or prophylactically to treat an animal having a disease or condition associated with caspase 6, such as Rieger's syndrome or ataxia telangiectasia, hyperproliferative disorder, a hamatopoietic disorder, a bone metabolism or cholesterol disorder, various types of cancer, neurological conditions such as Alzheimer's disease and other deregulated apoptotic pathological conditions. This polynucleotide sequence represents a mouse caspase 6 oligonucleotide relating to the invention. NOTE: This phosphorothicate oligonucleotide sequence has 2'-MOE wings and
                                                                                                                                                                                                                                              nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6 in cells or tissues. The oligonucleotides can be administered
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antilipaemic; osteopathic; caspase 6; Rieger's syndrome; bone metabolism; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder; haematopoietic disorder; cancer; neurological; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200229066-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an antisense oligonucleotide compound of 8 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       An antisense oligonucleotide of 8 inhibits caspase 6, is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Brown-Driver VL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-OCT-2000; 2000US-00679299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-OCT-2001; 2001WO-US030871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    apoptotic; mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse caspase 6 antisense inhibition related oligo SEQ ID No 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAL40381 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 91; 141pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; nootropic; neuroprotective; ophthalmological
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treating Rieger's syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to 50 nucleotides in length that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 6; Le
3.8e+04;
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Query Match

Sequence 20

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5 A; 6 C; 3 G; 6 T; 0 U; 0 Other;

Score

13;

DB

6; Length

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RESULT 49
AAL46719/c
ID AAL467
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AC AAL467
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DT 20-AUG
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DB Human
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KW Human;
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ABK70785/c
ID ABK7077
XX ABK7077
XX ABK707
XX Human
XX TSP1;
XW FG0189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a TSP1 (thrombospondin 1) domain-containing polypeptide comprising the proteins appearing as AAU80188 and AAU80189, encoded by cDNAs designated FG05969 and FG01896. Also included are proteins that are 50% homologous to the proteins and a polypeptide having at least one deletion, replacement, addition or insertion of amino acid in the proteins and having at least sepetitions of the TSP1 domain. The polypeptide can be use in drug compositions particularly for disorders associated with angiogenesis and vasculogenesis. The present sequence is a sequencing primer for the cDNAs of the invention
Human; serine-threonine protein kinase; cancer; diabetes; obesity; central nervous system disorder; inflammation; gene therapy; COPD;
                                                                             Human serine-threonine protein kinase coding sequence PCR primer #1
                                                                                                                                  20-AUG-2002
                                                                                                                                                                                       AAL46719
                                                                                                                                                                                                                                     AAL46719 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TSP1 domain-containing polypeptide useful for drug compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 15; 51pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-378268/41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK70785;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                  206 ACCTAGACCCTGG 218
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YOSHITOMI PHARM IND KK.
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thes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides the protein and coding sequences of a human serine-threonine protein kinase. The sequences can be used in the diagnosis, treatment and prevention of cancers (e.g. lenkaemia, lymphoma or melanoma), CNS disorders (e.g. Parkinson's disease, stroke, or traumatic brain injury), diabetes, eating disorders (e.g. obesity, anorexia, or cachexia), allergies, anaphylaxis, asthma, inflammation and chronic obstructive pulmonary disease (COPD). The present sequence is a PCR primer for the coding sequence of the invention
                                                                                                                                                                                                                                                                                                                              Human; ribonuclease L; antisense modulation; cytostatic; antimicrobial; antiinflammatory; antitumour; ribonuclease L expression inhibitor; antisense gene therapy; infection; aberrant apoptosis; cancer; tumour; inflammation; phosphorothioate; 2'-O-methoxyethyl; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human ribonuclease L antisense oligonucleotide SEQ ID NO:61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New human serine-threonine protein kinase and encoding polynucleotides, useful for diagnosing, treating and preventing central nervous system disorders (e.g. stroke), diabetes, or cancers (e.g. leukemia).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neuroprotective; antiparkinsonian; cerebroprotective; cytostatic; antidiabetic; antiallergic; antiasthmatic; antidepressant; anorectic; antiinflammatory; immunomodulator; chronic obstructive pulmonary disease;
                                                                                                                                                                                                                                                                            Homo sapiens
                            modified_base
                                                                                                                                                  modified_base
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13; Conserv
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Conservative
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Pred. No.
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                                                                                                                                                                                   RESULT 51
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a compound (I) of 8-50 nucleobases in length targeted to a nucleic acid molecule (II) encoding ribonuclease L (III), and which specifically hybridises with (II) and inhibits expression of (III), where (I) specifically hybridises with at least an nucleobase portion of an active site on (II). (I) has cytostatic, antiniflammatory and antitumour activities, and can be used as a ribonuclease L expression inhibitor and in antisense gene therapy. (I) is useful for inhibiting the expression of ribonuclease L icells or tissues, and for treating an animal having a disease condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 associated with ribonuclease L, e.g. infection, aberrant apoptosis or cancer. (I) is also useful for modulating the process of RNA-mediated interference (RNA) in a cell or animal. (I) is also useful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation. (I) is useful as a tool in differential and/or
                                                                                                                                                                                                                                                                                                                                                                                                 pathway, and in antisense gene therapy. The present sequence represents human ribonuclease L chimeric phosphorothioate antisense oligonucleotide which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                combinatorial analyses to elucidate expression patterns of a portion of the entire complement of genes expressed within cells and tissues. (I) also useful for research, therapeutics and diagnostics. (I) is also useful for distinguishing functions of various members of a biological
                                                                                                                                                  AAD53632 standard;
                                                                                                                                                                                                                                                                                                                                                                  Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; Page 78; 106pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel antisense compound which is targeted to nucleic acid encoding ribonuclease L, and inhibits expression of ribonuclease L protein, for treating diseases or conditions resulting from infections and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-313248/30
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Antisense; human; PTPN2; autoimmune

Human PTPN2 28-MAY-2003 AAD53632

antisense oligonucleotide, ISIS #135690.

protein tyrosine phosphatase non-receptor type 2; disorder; hyperproliferative condition; cancer;

(first entry

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ACH60223,
                                                                                                                                                              RESULT 52
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                                                                                                                                                                                                                                                                        Query Match
Best Local
Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotide compound, for diagnosing, preventing and/or treating conditions associated with aberrant expression or activity of protein tyrosine phosphatase non-receptor type 2 (PTPN2) e.g. cancer.
                                        DNA target sequence #9359 useful in array for genetic analyses
                                                                          17-OCT-2003
                                                                                                                               ACH60223 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to antisense compounds targetted to a nucleic acid a encoding human protein tyrosine phosphatase non-receptor type 2 (PTPN2). Antisense compounds of the invention are used for the diagnosis, prevention and treatment of diseases or conditions associated with PTPN2 such as autoimmune disorder, hyperproliferative condition e.g. cancer, or
                                                                                                                                                                                                                                                                                                                                                     such as autoimmune disorder, hyperproliferative condition e.g. cancer a haematopoietic disorder. The invention is useful in antisense gene therapy. The present sequence is an antisense oligonucleotide targett to human PTPN2 DNA. This oligo is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
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ilarity 100.0%;
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/note= "Phosphorothioate backbone; All cytidine residues
|rre 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                 7 T; 0 U; 0 Other;
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17-OCT-2003 ACH60616

(first

entry)

ACH60616 standard;

DNA;

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ВÞ

Gene expression analysis; array; hybridisation; genetic va tag-labelled compound; gene family; in situ hybridisation;

DNA target sequence #9752 useful in array for genetic analyses.

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5 77

AGACCTACCTGTG 89 AGACCTACCTGTG 3

Matches Query Match

Local

Similarity

100.0%; 2.2%;

Score 13; Pred. No.

DB 8; Le

Length 20 Indels

Conservative

0

Mismatches

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Comprises 9 or more consecutive nucleotides from at least one of 14936 comprises 9 or more consecutive nucleotides from at least one of 14936 comprises 9 or more consecutive nucleotides from at least one of 14936 comprises mismatch, antisense match or antisense mismatch oligonucleotides. The probes may be used in an array comprising at least 10 distinct conclude acid probes. The array is useful in monitoring gene expression levels by hybridisation to a DNA library, in analysing genetic consecutive, and in hybridising tag-labelled compounds. The probes are useful for identifying family members of a gene. The probes are also consecutive in in situ hybridisations, in screening CDNA or genomic libraries (or derived subclones) for additional clones containing segments of DNA contains the previously isolated and sequenced, in Southern, northern, and thave been previously isolated and sequenced, in Southern, northern, cor dot-blot hybridisation of genomic DNA to identify or detect the sequence of any gene or detect specific mutations in any gene, and in mapping the 5' termini of manA molecules by primer extensions. The compling the southern of the invention are also useful as PCR primers. The invention provides a large collection of nucleic acid sequences complementary to particular genes with a wide range of analytical uses. ACH50865-ACH65260 represent the target sequences of the invention. Note: The sequence data for this patent was obtained in electronic format contains and the proper sequences of the invention. Note: The sequence data for this patent was obtained in electronic format the target sequences of the invention. Note: The sequence data for this patent was obtained in electronic format the sequence of the invention of the constant was obtained in electronic format the sequence of the invention of the constant of the constant of the sequence of the invention.
Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to nucleic acid sequences that are complementary to particular genes, and can be used as probes for a variety of analyses such as gene expression analysis. Each probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New probe array useful e.g. for monitoring gene expression levels, for analyzing genetic variations, or for hybridizing tag-labeled compounds, comprises multiple nucleic acid probes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 9359;
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Gene expression analysis; array; hybridisation; genetic variation, tag-labelled compound; gene family; in situ hybridisation;
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                                                          DNA target sequence #10087 useful in array for genetic analyses.
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                                                                                                                                                                                      ACH60951 standard; DNA; 20
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RESULT 55 ACH60952

ACH60952 standard; DNA; 20

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ACH60952

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AGACCTACCTGTG 89 AGACCTACCTGTG

Matches Query Match Best

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Similarity

2.2%;

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Conservative

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entry)

DNA target 17-OCT-2003

Gene expression analysis; array; hybridisation; genetic variation, tag-labelled compound; gene family; in situ hybridisation;

sequence #10088 useful in array for genetic analyses.

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CC variations, and in hybridising tag-labelied compounds. The probes are cuseful for identifying family members of a gene. The probes are also cuseful in in situ hybridisations, in screening cDNA or genomic libraries conserved in the probes are also cuseful in in situ hybridisations, in screening cDNA or genomic libraries cor derived subclones) for additional clones containing segments of DNA conditions that have been previously isolated and sequenced, in Southern, northern, cor detect hybridisation of genomic DNA to identify or detect the conditions of any gene or detect specific mutations in any gene, and in mapping the 5' termini of mRNA molecules by primer extensions. The cucleic acid sequences of the invention are also useful as PCR primers. The invention provides a large collection of nucleic acid sequences complementary to particular genes with a wide range of analytical uses. ACH50865-ACH65260 represent the target sequences of the invention. Note: The sequence data for this patent was obtained in electronic format conditions are also useful as PCR primers.
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20 BP; 3 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
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RESULT 56
ACH60727
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Gene expression analysis; array; hybridisation; genetic vatag-labelled compound; gene family; in situ hybridisation;
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                                                             DNA target sequence #9863 useful in array for genetic analyses
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17-OCT-2003 ACH60447;

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ACH60447 standard; DNA; 20

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Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation; DNA target sequence #9583 useful in array for genetic analyses

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                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
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Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation;
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                                                           DNA target sequence #9584 useful in array for genetic analyses.
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ACH60451

ACH60451 standard; DNA; 20

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17-OCT-2003

(first entry

Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation;

DNA target sequence #9587 useful in array for genetic analyses

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AGACCTACCTGTG

Matches Query Match

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l Similarity
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DB 8; Le 3.8e+04;

Length 20

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AGACCTACCTGTG 89

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The probes may be used in an array comprising at least 10 distinct connected probes. The array is useful in monitoring gene expression converted by hybridisation to a DNA library, in analysing genetic converted by hybridisation to a DNA library, in analysing genetic converted in an analysing genetic converted by hybridisations and in hybridisations, and in hybridisations, in screening cDNA or genomic libraries converted subclones) for additional clones containing segments of DNA that have been previously isolated and sequenced, in Southern, northern, correctly for a detect specific mutations in any gene, and in mapping the 5° termini of mRNA molecules by primer extensions. The mucleic acid sequences of the invention are also useful as PCR primers. CC complementary to particular genes with a wide range of analytical uses. CC ACH50865-ACH65260 represent the target sequences of the invention. Note: CC directly from the USPTO web site at sequence of nectronic format condirectly from the USPTO web site at sequence.
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20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                               RESULT 60
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                                                                                                                                                                                                                                                                                                                                                                            Matches
Gene expression analysis; array; hybridisation; genetic variation, tag-labelled compound; gene family; in situ hybridisation;
                                                             DNA target sequence #10091 useful in array for genetic analyses.
                                                                                                                                                                                       ACH60955 standard; DNA; 20
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llarity 100.0%;
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RESULT 61
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3.8e+04;
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Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation; DNA target sequence #9247 useful in array for genetic analyses

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antisense oligonucleotide; human; mdm2; hyperproliferation;
hyperproliferative disorder; cancer; psoriasis; fibrosis;
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Pred. No.
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hes 0;
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07-APR-2000;

2000DE-01019173 2001WO-IB000713.

06-APR-2001; 18-OCT-2001 WO200177384-A2 Homo sapiens central nervous Cligonucleotide primer SEQ ID NO 334825 for detecting SNP TSC0038427.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;

system; gastrointestinal;

respiratory;

immune; metabolic

cancer; CNS;

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ABI34852/c
ID ABI348
XX
AC ABI348
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AC ABI348
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SNP; S
W SNP; S
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antisense compound targeted to 5' untranslated region, region, or intron:exon junction of nucleic acid molecule encuseful for treating e.g. cancer, psoriasis or restenosis by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABI34852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABI34852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 0 A; 10 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention comprises antisense oligonucleotides which are targeted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 9; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mdm2 expression.
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Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-2001; 2001US-00005344.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 atherosclerosis; restenosis; apoptosis modulation; p21;
2'-methoxyethoxy-residue; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2003-577263/54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAGCCCGAGGGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ID NO 41; 289pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 9; Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.8e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  jion, coding
encoding mdm2,
s by inhibiting
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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RESULT 64
ABI35151
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99989, ABF00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989, and ABI00010-ABF89989, and the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the interval of the printed specification of the printed specification.
                                                                                                                                                                olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; peptide nucleic acid; cytosine methylation; cardiovasc, central nervous system; gastrointestinal; respiratory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide primer SEQ ID NO 335124 for detecting SNP TSC0038615.
                   Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin
                                                                                                             WPI; 2001-657177/75
                                                                                                                                                                                                                                                                           07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABI35151 standard;
methylation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 334825; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                      (EPIG-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAGATTATTTT
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  status
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No.
                                                                                                                                                                Mismatches
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. 1.2e+05;
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Claim 1; SEQ ID NO 335124; 29pp + Sequence Listing; German.

describes novel oligonucleotide primers or peptide nucleic

Sequence

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ស្ត្រាស្ត្រស្ត្រាស្ត្រស្ត្រស្ត្
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Best Local
                                                                       acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0001-ABC99989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metaboli
                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 360284; 29pp + Sequence Listing; German
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                                                   ftp.wipo.int/pub/published_
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAAGATTATTTT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12;
Pred. No.
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Query Match Best Local Matches

Similarity

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Score 12; DB 5; I Pred. No. 1.2e+05;

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Matches 12
                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation, ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                Sequence 12 BP; 1 A; 0 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 296679; 29pp + Sequence Listing; German.
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                          301 AACCCCAACCTC 312
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                                                                         Similarity
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                                                        Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer SEQ ID NO 296679 for detecting SNP TSC0017210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                system;
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                                                                       2.0%; Score 12; DB 5;
100.0%; Pred. No. 1.2e+0
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RESULT 67 ABI78279 ID ABI78

ABI78279 standard; DNA; 12

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Homo sapiens

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ABC01330
ID ABC01330
XX ABC01
AC ABC01
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                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9988, ABF00010-ABF99889, ABH00010-ABH9988 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                         Oligonucleotide SEQ ID NO 1321 for detecting
                                                                                                                                                                                                                                                                                                                                                                                               ABC01330 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 12 BP; 4 A; 0 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-FEB-2002
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           WPI; 2001-657177/75
                                                             Olek A,
                                                                                                                                                                       07-APR-2000;
                                                                                                                                                                                                                         06-APR-2001; 2001WO-IB000713.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                  EPIGENOMICS
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                                                             Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; DNA;
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                                                                                                                                                                       2000DE-01019173
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Best Local S
Matches 12
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                                                                                                                                                                                          Claim 1;
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Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Claim 1; SEQ ID NO 54269; 29pp + Sequence Listing; German

acid (PNA) oligomers for detecting single nucleotide polymorphisms (S) and cytosine methylation status in chemically pretreated genomic DNA. oligonucleotides are used for diagnosis and/or prognosis of cancer and ftp.wipo.int/pub/published_pct_sequences describes novel oligonucleotide primers or peptide nucleic NOTE: The sequence cancer and a respiratory, ABC00010 (SNP) but

Sequence 13 BP; 8 A; 0 C; 1 G; 4 T; 0 U; 0 Other;

0 Score 12; Pred. No. Mismatches DB 5; Le 1.2e+05; Length 13 Indels 0 Gaps 0

쁌

Oligonucleotide SEQ ID NO 195374 for detecting SNP TSC0048069.

central nervous system; gastrointestinal; respiratory; immune; metabolic single nucleotide polymorphism; human; diagnosis; PNA; ide nucleic acid; cytosine methylation; cardiovascular; primer; 88

Berlin <u>.</u>

Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine 18

SEQ ID NO 195374; 29pp + Sequence Listing; German

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The

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ABF99192
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Best Local Similarity
Matches 12; Conserv
                    Query Match
Best Local !
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                                                                                             range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABH00010-ABH82073 represent the oligomers described in the Invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligomers are also used for detecting cell type differentiation. ABC0001-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF92073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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                                                                  Sequence 13
                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 199189; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                         methylation status.
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ilarity 100.0%;
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2.0%; Score 12; DB ilarity 100.0%; Pred. No. 1.2 Conservative 0; Mismatches
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Pred. No.
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Indels

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22-FEB-2002 ABH06779 ABH06779

(first entry)

standard;

DNA; 13

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RESULT 72
ABH06778/c
ID ABH06778/c
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XX Oligon
XX SNP; se
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RESULT 73
ABH06779
ID ABH06
XX
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XX SNP; E
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DMA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABC99889, ABF00010-ABC99889, ABF00010-ABC99889, and MBI00010-ABC99889, and Colligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;
                                                                                                                                                                                          Oligonucleotide SEQ ID NO 195373 for detecting SNP TSC0048069.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 13
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                                                                                                                                                                                                                                                         ABF95376
                                   18-OCT-2001.
                                                                 WO200177384-A2
                                                                                                                                                                                                                                                                                       ABF95376 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide SEQ ID NO 206756 for detecting SNP TSC0050584
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 7 A; 2 C;
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                                                                                                                             system;
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06-APR-2001; 2001WO-IB000713.

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ID ABF88
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ID ABF88
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                         Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligomers are also used for detecting cell type differentiation. ABC000 -ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, bu was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                               EPIGENOMICS
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                                                                                                                                                                                                                Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; DNA; 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTATTTTTACT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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Pred. No.
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RESULT 76
ABF35048/c
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Best Local
Set of oligonuclectides, useful for diagnosis and cell typing, is designed to detect single-nuclectide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 135045; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; ide nucleic acid; cytosine methylation; cardiovascular; primer; ss; ral nervous system; gastrointestinal; respiratory; immune; metabolic
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Pred. No.
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12

TTTTGTAGAAAA 461

Matches Best Query Match

POCST

Similarity

2.0%;

Score 12; DB 5; L Pred. No. 1.2e+05;

Length 13; Indels

Conservative

0

Mismatches

°.

Gaps

0

Sequence

13

BP; 5

A; 2 C; 0 G; 6 T; 0 U; 0 Other;

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RESULT 77
ABP51823/c
ID Oligon
XX SNP; g
XW peptid
XW centra
XX W02001
XX WPI; G
PT Olek ;
XX WPI; G
ID ABP1
XX WPI; G
ID ABC91
CC Cantra
XX Contra
CC Cantra
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                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99889, ABH00010-ABF99899, ABH00010-ABF9989989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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Pred. No. 1.2e+05;
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                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide SEQ ID NO 54270 for detecting SNP TSC0014903.
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                                                                                                               Claim 1; SEQ ID NO 54270; 29pp + Sequence Listing;
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This invention describes acid (PNA) oligomers for and cytosine methylation

novel oligonucleotide primers or peptide nucleic detecting single nucleotide polymorphisms (SNP) status in chemically pretreated genomic DNA. The

Query Match

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                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                              Seguence 13
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ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal, respirators.
                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                      Oligonucleotide SEQ ID NO 199190 for
                                                                                                                                                                                                    ABF99193;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                     22-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13 BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  was obtained in electronic format from W ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 113244; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     designed to detect single-nucleotide methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

Claim 1; SEQ ID NO 151819; 29pp + Sequence Listing;

German

Set of oligonucleotides, designed to detect single

tides, useful for diagnosis and cell single-nucleotide polymorphisms and

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07-APR-2000; 2000DE-01019173 06-APR-2001; 2001WO-IB000713

EPIGENOMICS Piepenbrock C,

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18-OCT-2001 WO200177384-A2 Homo sapiens

central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC-ABC99989, ABFC0010-ABF99989, ABHC00010-ABF99989 and ABI00010-ABF89989

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RESULT 88
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Matches 12; Conserv
                                                                                                         range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABB99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  represent the oligomers described in the invention. No data for this patent did not form part of the printed was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
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                                                                Sequence
                                                                                                                                                                                                                                                                                               Claim 1;
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RESULT 90
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AC ABF36
AC AB

ABF35049 standard;

DNA; 13

βP

21-FEB-2002 ABF35049

(first entry)

Oligonucleotide SEQ

ID NO

135046

for

detecting

SNP TSC0033667

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                               Matches
                                               Query Match
Best Local
                                                                                                                                       acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin methylation status.
                                                                                            Sequence 13 BP; 1 A; 0 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                      was obtained in electronic format from W ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 262079; 29pp + Sequence Listing; German.
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301 AACCCCAACCTC 312
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12; Conserv
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                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
                                                                                                                                                                                                               Oligonucleotide SEQ ID NO 262080 for detecting
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07-APR-2000;
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                                                                                         WO200177384-A2
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2000DE-01019173
                                                                                                                                                                                                                                             (first entry)
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Disclosure; Page 30;

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                     New promoter sequences for causing expression of a structural gene especially agronomic gene or open reading frame in a plant cell, comprises engineered versions of the maize ubiquitin promoter.
                                                                                                                                                                                                       09-JUN-2000; 2000US-00590558.
                                                                                                                                                                                                                                  08-JUN-2001; 2001WO-US018689
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                                                                                                                                                                                                                                                                                                                         Arabidopsis thaliana.
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                                                                                                                 WPI; 2002-122117/16.
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                                                                                                                                                                           (PROD-) PRODIGENE INC
                                                                                                                                                                                                                                                                                                                                                     expression; maize; ubiquitin promoter; Ubi-1; HSE; shock element; agronomic gene; HY5 transcription factor;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                      Single stranded DNA molecule, which binds adenosine or adenosine-5'-phosphate - useful as purification reagent, or for determination of adenosine tri:phosphate subcellular localisation in vivo.
                                                                                                                                                                                                                                                                                                                                       Szostak JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Adenosine; adenosine-5'-phosphate; adenosine triphosphate; ATP; ligand; purification; reagent; isolation; determination; subcellular localisation; catalyst; assay; SELEX; Systematic Evolution of Ligands by EXponential enrichment; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT79478 standard; DNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 2 A; 5 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                19-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5631146-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ribulose-1,5-biphosphate carboxylase gene, used in the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12; Conserv
                                                                                                                                                                                                                                                                                                                                                                            GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGCCACGTGGAA 174
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                       Huizenga DE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 adenosine or adenosine 5'-phosphate
                                                                                                                                                                                                                                                                                                                                                                                                              95US-00375116
                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-00375116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 6; Lo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             binding;
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binding single stranded DNA molecule, which can be used as a purification reagent for the isolation of adenosine or an A5P, or to determine the subcellular localisation of, e.g. adenosine triphosphate (ATP), in vivo. The DNA molecule was prepared by contacting DNA molecules having a region of random sequence with adenosine or A5P (preferably ATP), isolating a subpopulation by partioning DNA molecules which specifically bind the adenosine or A5P, amplifying the subpopulation in vitro and repeating the process 4 times to obtain a single stranded DNA molecule capable of

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CTCCTGGAGCCC

14

The present sequence is an adenosine or adenosine-5'-phosphate (A5P)

Claim 3; Col 63-64; 55pp; English.

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RESULT 94
AAQ40305
ID AAQ40
XX AAQ40
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                                                           Query Match
Best Local S
Matches 12
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Best Local
                                                                                                                                                                                                        a novel DNA construct under control of transcriptional initiation region operative in plants. The construct is used to produce plants in which GPDH expression is modified; in partic. the constructs produce antisense RNA to down regulate expression. The transcriptional initiation region is a constitutive promoter. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         binding adenosine or A5P, i.e. Systematic Evolution of Ligands by EXponential enrichment (SELEX). Catalytic DNA produced using the method can be used as in vitro or in vivo catalysts, or to detect the presence of the ligand. They may also be used in assays to detect molecules modified by the DNA, which are not themselves ligands, e.g. DNA phosphorylated by a polynucleotide kinase catalyst. The DNA molecule has significant advantages over ligand binding and catalytic RNA in terms of
                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-OCT-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence of PCR primer oligo lambda-F for amplification of a fragment pTOM38 encoding glyceraldehyde-3-phosphate dehydrogenase (GPDH).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ40305;
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                                                                                                                                                                                                                                                                                                                                                                                          Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             modify ripening characteristics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA constructs for anti:sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-APR-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Glyceraldehyde-3-phosphate dehydrogenase; inhibitor; antisense RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
10-AUG-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ40305
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12; Conserv
                                                           l Similarity
12; Conserv
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                                                                                                                                                                                                                                                                                                                                       sequence is
                                                                                                                                                                                                                                                                                                                                                                                     Fig 2; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; cDNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTTCCTCCGCTA 191
               CTCCTGGAGCCC
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                                                                                                                                                               BP;
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                                                                                  2.0%;
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                                                                                                                                                               6 C;
                                                                                                                                                                                                                                                                                                                                       a fragment of cDNA for tomato cytosolic GPDH used
               257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IND PLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ray JA,
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                                                                                                                                                               G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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                                                                                       Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                inhibition of GPDH in plants - used to
                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schuch WW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2; L,
1.2e+05;
                                                                                                              DB 2;
                                                                                     1.2e+05;
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AAQ89599
ID AAQ89599
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XX Kappa
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RESULT 96
AATT32436
ID AATT32
XX AATT32
XX AATT32
XX 30-SE
XX 30-SE
XX Wasp;
KW Wasp;
KW Lepid
XX Drime
XX Synth
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Best Local :
                                                                                                                                                   Wasp; venom;
Lepidoptera;
primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A commercial cDNA library prepd. in lambda gtll from mRNA obtd. from human breast tissue removed during the third trimester of pregnancy was screened with rabbit anti-bovine kappa-casein cDNA. The cDNA insert of a recombinant phage was amplified by PCR using the primer given in AAQ89595 (located 13-27 bp upstream of the EcoRI site of lambda gtll) and AAQ896 (located 8-22 bp downstream of the EcoRI site)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ89599 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA encoding human kappa-casein - used for the prodn. of large amts. highly purified kappa-casein milk protein for infant use.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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     30-MAY-1996
                                                                                                      Synthetic.
                                                                                                                                                                                                                                                      PCR primer lambda gtll forward.
                                                                                                                                                                                                                                                                                                           30-SEP-1996
                                                                                                                                                                                                                                                                                                                                                          AAT32436;
                                                                                                                                                                                                                                                                                                                                                                                                            AAT32436 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTCCTGGAGCCC 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTCCTGGAGCCC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jeffers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Col 11; 14pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         larity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                          neurotoxin; insecticide; biological
insect; Bracon hebetor; polymerase 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
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                                                                                                                                                                                                                                                                                                                                                                                                            DNA; 15 BP
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.2e+05;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 2; Length 15;
                                                                                                                                                                             chain reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                               PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sert of a
n AAQ89599
1 AAQ89600
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RESULT 97
AAX56935/c
ID AAX569
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Best Local Similarity
Matches 12; Conserv
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19-JAN-1995;
29-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The PCR primer pair lambda gtl1 forward (AAT32436) and lambda gtl1 reverse (AAT32437) were used to screen for the presence of a cDNA insert in plaque-purified phage obtd. from a Bracon hebetor wasp cDNA library. The primers are specific for phage lambda gtl1 and flank the BcoRI cloning site. cDNA clone pBrhTX-1(a)1.1 (AAT32438) was identified that codes for the toxin (a) subunit (AAR99568) of the wasp neurotoxin BrhTX-1. This can be utilised in breeding of biological control agents used to
                                                                                                                                                                                                                                                                    HIV-1 proviral DNA fragment 18.
                                                                                                                                                                                                                                                                                                                                                  AAX56935 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-NOV-1995;
        Derivatised solid supports and reagents for oligo:nucleotide synthesis and new oligo:nucleotide phosphoramidate conjugates.
                                                                                                                       13-MAY-1994;
                                                                                                                                                                                                                                  DNA-targeting conjugate; viral DNA-binding agent;
                                                                                                                                                                                                                                                                                           16-OCT-2003
15-JUL-1999
                                                                                                                                                                                                                                                                                                                            AAX56935;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 20; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bracon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Windass JD,
                                           WPI; 1996-010846/01.
                                                                 Watanabe
                                                                                       (SLOK )
                                                                                                                                                                    23-NOV-1995.
                                                                                                                                                                                         WO9531434-A1
                                                                                                                                                                                                              Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        combat insect pests
                                                                                                                                             12-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ZENE ) ZENECA LTD.
(CSIR ) COMMONWEALTH SCI & IND RES
                                                                                                                                                                                                                                                                                                                                                                                                                               246
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hebetor toxins and DNA encoding lagents to combat insect pests.
                                                                                      SLOAN KETTERING INST CANCER RES ZW BIOMEDICAL RES AG.
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                                                                                                                                                                                                                                                                                                                                                                                                                              CTCCTGGAGCCC 257
                                                                                                                                                                                                                                                                                                                                                                                                          CTCCTGGAGCCC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Duncan
                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                               Ren W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94GB-00023540.
95GB-00001074.
95GB-00013293.
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                                                                                                                       94US-00242664.
                                                                                                                                             95WO-US006379
                                                                                                                                                                                                                                                                                                                                                  DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       쮼,
                                                                 Weil
                                                                                                                                                                                                                                  anticancer drug; viral DNA-cleaving agent;
solid support; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Baule VJ,
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                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Christian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2; Lo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            them
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RESULT 98
AAT14822
ID AAT14
R3CH2CH2CH2CH2R4, where: R3 = a protecting group; and R4 = OH or an H-
C phosphonate, alkanephosphonate, phosphortiester, phosphite triester,
c phosphite diester, phosphoratioate, phosphoramidate, phosphoramidate,
c phosphite diester, phosphoratioate, phosphoramidate,
c propagating an oligonucleotide 5', phosphate, a process for
c preparing a solid support useful for preparation of an oligonuclectide 3'
c phosphate, a process for preparing an oligonucleotide 3', 5'-diphosphate and a
c process for preparing an oligonucleotide 3', 5'-diphosphate. The
c process for preparing an oligonucleotide 3', 5'-diphosphate. The
c process for preparing an oligonucleotide 3', 5'-diphosphate. The
c targeting conjugates, e.g. with anticancer drugs or viral (e.g. HIV) DNA-
c cleaving or -binding agents. The process for preparing oligonucleotide
3', 5'-diphosphates is simple and suitable for use in automatic DNA
c synthesisers. This sequence represents a fragment of the HIV-1 provirus
genome, used to describe the method of the invention. (Updated on 16-OCT-
2003 to standardise OS field)
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel derivatised solid supports of formula S'-L-Z-CH2CH2-R, where: S' = a solid support; L = a bond or an (in)organic linker; Z = SO2 or S-S, R = OH, an H-phosphonate alkanephosphonate, phosphotriester, phosphite triester, phosphorate diester, phosphorothioate, phosphorodithioate, phosphoramidate or phosphoramidite group, OR1, SR1, an optionally substituted or modified nucleotide (N'), or an oligonucleotide of formula (N')gR2; g = 1-200; R1 = a protecting group; R2 = an H-phosphonate, alkanephosphonate, phosphotriester, phosphite triester, phosphoramidate, alkanephosphonate, phosphorodithioate, phosphoramidate or phosphoramidite group, OH, OR1, SR1 or optionally substituted alkanephosphonate, phosphorodithioate, phosphoramidate or phosphoramidite group, OH, OR1, SR1 or optionally CH2CH2CH2CHOCH2CH2CH2CHOCH2CH2CH2CHCRILA Also mentioned are compounds of formula
                 DNA encoding histiocyte-secreted factor and its variants - anti-tumour agent and for studying tumour regression, havir cytotoxicity compared to TNF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Histiocyte-secreted factor; HSF; cytokine; antitumour; tumour; therapy; polymerase chain reaction; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                   WPI; 1996-239499/24
                                                                                                                                                                                                                                                                26-OCT-1994;
                                                                                                                                                                                                                                                                                                                26-0CT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                               WO9613586-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lambda gt11 flanking sequence 5′ primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          175
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TIGCICITCCIC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TIGCICTICCIC 186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 9 A; 1 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                94JP-00297780
                                                                                                                                                                                                                                                                                                                95WO-JP002200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12; DB 2; Leng; Pred. No. 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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                                                                   useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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RESULT 99
AAT35227
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                            Matches
                                           Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lambda gtl1 5' (AAT14822) and 3' (AAT14823) primers were used in a lambda gtl1 cDNA insert screening kit to identify insert DNA in clones obtd. by the PCR amplification of human histicoytic lymphoma DNA from a lambda gtl1 library. A genomic clone (AAT14818) coding for human HSF (AAR96800).
                                                                                                                     PCR primers ZC2682 (AAT35226) and ZC2683 (AAT35227) anneal 5' and 3' to the ECRI cloning site of lambda gtll. They were utilised in the ECRI amplification of cDNA inserts in clones that had been obtd. from a human placenta cDNA library in lambda gtll using a partial clone as probe (see also AAT35222-25). 2 Different PCR products were generated (AAT35220-21) that respectively coded for cytoplasmic antiproteinases 2 and 3 (AAR99253
                                                                                                                                                                                                                                             Human cytoplasmic antiproteinase-2 (CAP-2) and CAP-3 - serine protease inhibitors useful in the purification of proteins and in the treatment inflammatory diseases and apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytoplasmic antiproteinase-2 protein; CAP-2; CAP-3; serpin; serine protease inhibitor; antiinflammatory; apoptosis; polymerase chain reaction; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT35227 standard;
                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                Sprecher
                                                                                                                                                                                                                                                                                                                                                                                      08-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT35227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                     Example 1; Page 43; 50pp; English
                                                                                                                                                                                                                                                                                                     WPI; 1996-393014/39.
                                                                                                                                                                                                                                                                                                                                                                                                               02-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                           15-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytoplasmic antiproteinase PCR primer ZC2683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; Page 30; 52pp; English
                                                                                                                                                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       novel cytokine, was identified
  246
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                            l Similarity
12; Conserv
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 CICCIGGAGCCC 257
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                                                                                BP; 2
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                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                 96WO-US001288
                                                                                A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA; 15
                                       2.0%;
                                                                                6 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                            <u>.</u>
                                                                               G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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Pred. No.
                                       Score 12; DB 2; L
Pred. No. 1.2e+05;
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                             Mismatches
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RESULT 100

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                                                                                                                           subunit cDNA of the invention. The specification of an insecticidal toxin subunit cDNA of the invention. The specification provides a 1811 base pair spliced RNA (AAV17183) derived from a Bracon hebetor genomic clone that encodes at least two of the insecticidal toxin subunits shown in sequences AAW52124-W52128. The spliced RNA can hybridise with extension products prepared from a 564 base pair (AAV17145) template with 6 specified primers. A nucleic acid encoding at least one of the specified subunits can be modified so that mRNA instability motifs and/or fortuitous splice sites are removed, or insect-pest preference codons are used, so that expression of this nucleic acid in insect cells yields practically the same protein as unmodified nucleic acid in its endogenous companism. The nucleic acid encoding an insecticidal toxin subunit can be complementary to a sequence that hybridises under specified conditions to these nucleic acids, organisms regenerated from these cells and pathogens containing these nucleic acids and insecticidal compositions comprising a combinations of the toxin subunits are all used for control of insects. The nucleic acids are used to produce recombinant baculoviruses for
                                              Query Match
Best Local :
                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-MAY-1996;
22-MAY-1996;
22-MAY-1996;
22-MAY-1996;
22-MAY-1996;
22-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid encoding a combination of insecticidal sub:unit(s) of wasp toxin - and related transformed cells, insect pathogens and
                                                                                                                                                                                                                                                                                                                                                                                                                                      combinations of proteins, useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-018430/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Duncan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                insecticidal toxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Insecticidal toxin subunit cDNA amplifying primer 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-JUN-1998
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                                                                                      Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 61; 84pp; English
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  246
                               12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZENECA LTD.
COMMONWEALTH
                                            Similarity
  CTCCTGGAGCCC
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                                                                                        BP; 2 A;
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                               Conservative
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96GB-00010738.
96GB-00010739.
96GB-00010748.
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                                            2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                      6 C; 5 G; 2 T; 0 U; 0 Other;
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                                           Score 12; DB 2; Li
Pred. No. 1.2e+05;
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AAX61290
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Matches
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                   Mosquito; Aedes aegypti; IgB-binding protein; immunoglobulin; bite; inflammatory reaction; allergen; skin test; immunotherapy; rash; erythema; oedema; induration; pain; itch; fever; systemic reaction; antigen; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is a primer for the cDNA encoding the recombinant mosquito salivary allergen rAed a 3. rAed a 3 can be used to determine sensitivity to mosquito bites by epi-cutaneous or intradermal testing, a substrate to which IgE/G bind (e.g. for mosquito allergy diagnosis) a substrate tresults of the skin tests) for immunotherapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Recombinant; mosquito; salivary allergen; rAed a 3; determination; bite sensitivity; epi-cutaneous test; skin test; intradermal test; allergy diagnosis; immunotherapy; desensitisation; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Recombinant mosquito salivary allergens for use in skin tests sensitivity - also as substrate for assaying allergen-specific immunoglobulin and for de-sensitisation immuno-therapy.
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                                                                                                                                                                                                             28-JUL-1999
                                                                                                                                                                                                                                                                                                                      AAX61290 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 4; Page 43; 82pp; English.
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                                                                                                                                                         Aedes aegypti saliva cDNA PCR primer SEQ ID NO:2.
                                                                                                                                                                                                                                                                 AAX61290
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Pred. No. 1.2e+05;
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RESULT 103
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ID AAZ397
XX AAZ397
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                     Caspase; serpin; inflammation; apoptosis; lung disease; human; CAP; neurodegenerative disease; heart; liver tissue; Alzheimer's disease; Parkinson's disease; amyotrophic lateral sclerosis; injury; trauma; hypoxy-ischaemia; cytoplasmic antiproteinase protein; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bites. The mosquito extract consists essentially of these antigens to allergens in mosquito saliva. The assays are applicable to patients presenting with rashes and other symptoms after mosquito bites, especially those with erythema, oedema and induration, pain or itch at the site(s) of mosquito bite(s), with or without fever. These patients are at risk for severe localized inflammatory reactions and systemic reactions to mosquito bites. The present sequence represents a PCR primer for an Aed a 3 cDNA clone AA22, which was isolated from Aedes aegypti (mosquito) saliva in an example from the present invention for the isolation of a cDNA encoding a 30 kDa IgE-binding protein
                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                              Human CAP DNA specific primer ZC2683.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15
       04-MAY-1998;
                                                   27-APR-1999;
                                                                                                   11-NOV-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ39756 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a mosquito extract consisting essential of antigens related solely to allergens in mosquito saliva. The isolate and purified recombinant mosquito salivary antigens are for use in skin
                                                                                                                                                  WO9957273-A2
                                                                                                                                                                                                                                                                   nootropic; neuroprotective; vasotropic; tranquilizer; vulnery; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-347473/29.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mosquito extract consisting of antigens to allergens in mosquito saliva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYMA-) UNIV MANITOBA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunoassays and immunotherapy for allergic reactions to mosquito
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12;
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       98US-00072275
                                                   99WO-US008949
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Pred. No.
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  RESULT 104
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The invention provides a method for treating a disease mediated by a caspase in an individual. The method comprises: administering a composition comprising a gene coding for an intracellular mammalian serpin in an amount sufficient to inhibit activity of the caspase upon transient expression of the gene in a target tissue affected by the disease, where the disease or the symptoms are treated. The method can be used for decreasing inflammation, for modulating apoptosis, for treating a lung disease, and for treating a neurodegenerative disease. The inflammation and apoptosis that can be treated are particularly in heart or liver tissue. It can be used for treating Alzheimer's disease, Parkinson's disease, amyotrophic lateral sclerosis, and acute injury such as a position for the name. Sequences A23975-56 represent PCR
                                                                                                                                                        primers specific for the human cytoplasmic antiproteinase protein (CAP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sprecher CA,
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                                                                            BP; 2 A; 6 C; 5
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                                                                        G; 2 T; 0 U; 0 Other;
    DB 3;
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Length 15
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12; Conserv
CTCCTGGAGCCC
                CTCCTGGAGCCC 257
                                 Conservative
                                        2.0%;
                                 0
                                        Score 12;
Pred. No.
                                 Mismatches
                                 1.2e+05;
hes 0;
                                 Indels
                                0
                                Gaps
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13-SEP-2001
                                                      AAH21253 standard; DNA; 15
(first entry)
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Kv4.1 screening amplimer SEQ ID 10

Human; Kv4.1; potassium channel protein; Kv4.2; autism; epilepsy; neurodegenerative disease; ischemia; stroke; Alzheimer's disease; Parkinson's disease; Huntington's disease; cardiac arrhythmia; memory; learning capacity; protein kinase activator; anti-arrhythmic; primer; SS

Homo sapiens.

DE19963612-A1

12-JUL-2001

29-DEC-1999; 99DE-01063612.

29-DEC-1999; 99DE-01063612.

(GENI-) FORSCHUNGSGESELLSCHAFT GENION MBH

WPI; 2001-426637/46.

testing potential pharmaceuticals, New potassium chammel subunit proteins, e.g. useful for identifying and anti-arrhythmic or neurological

2; Page 26; 50pp; German

This invention describes a novel potassium channel protein (I) that įs

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Query Match
Best Local S
Matches 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               channels containing (I) are used to identify and test: (i) compounds for treatment of neurodegenerative diseases (autism, epilepsy, ischemia, stroke; Alzheimer's, Parkinson's and Huntington's diseases) or cardiac arrhythmia, or those that improve learning capacity and memory; and (ii) activators of protein kinases. Host cells that express (I) can identify agents that do not interact significantly with channels and control It (a quickly activated transient current), so lack the side effects of known anti-arrhythmic agents. They also eliminate, or reduce, the need
The present invention relates to a method for ameliorating the effects skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor. IGF binding protein [IGFBP]-2 or IGFBP3, which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to despit the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                            Ameliorating the effects of a disorder, e.g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JUN-1999;
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1.2e+(
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                skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor. IGF binding protein [IGFBP]-2 or IGFBP], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense fastion. The method is useful for amaliorating the effects of psoriasis ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition of the retin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1, pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; aclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inhibits or reduces growth
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Best Local :
                                          inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAR45151 and AAR45153-745161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warrs, benign growths, cancers of the skin, a hyperneovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
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vessels c
                                                                                                                                                                                   The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Pactor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1, pityriasis; IGF binding protein; IGF9D-2; IGFBD3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                              Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
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or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGCCCGAGGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15 BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGCCCGAGGGGC 4
                                                                                                                                                                                                                                                                                        Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                             other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0140345P
                                                                                                                                                                                                                                                                                        67;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                   201pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Edmondson SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       쁌
                                                                                                                                                                                                                                                                                      English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15;
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Query Match Best Local

Sequence

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BP;

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7 C; 4 G; 3 T; 0 U; 0 Other;

Matches

1 Similarity 12; Conserv

Conservative 0;

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Score 12; Pred. No. Mismatches

DB 4; I

Length 15

Indels

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Gaps

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RESULT 108
AAP50005/c
ID AAP5000
XX AAP5000
XX AAP5000
XX AAP5000
XX AAP5000
XX ATTISE
KW GYTOST
KW GYTOST
KW GYTOST
KW HOTOST
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Best Local :
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                                                                                                                                                          skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor. IGF binding protein [IGFBP]-2 or IGFBP], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the facts of psoriasis, 1910 and present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition of the retina, the province of the skin, a service of the skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrheea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF50005;
                                                                                                  brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a method for ameliorating the effects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 8; Page 67; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wraight CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JUN-2000; 2000WO-AU000693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF50005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IGF-I oligonucleotide #965
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                365
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12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGCCCGAGGGGC 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AGCCCGAGGGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0140345P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GA,
                                                                  hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of.
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AGCCCGAGGGGC 376

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RESULT 109
ACD26289
ID ACD262
RESULT 110
AAQG6708/c
ID AAQG677
AX AAQG677
AC AAQG67
XX
DT 22-DEC
XX
XX
DE Primer
XX
KW HHV6;
KW polyme
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                                                                                                                                                                                                                                                                                                         В
                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                             The invention relates to a gene chip which is composed of a chip substrate and oligomeric nucleic acid probes fixed on the substrate. The probe is composed of a detecting region and a stem region, where a hairpin-shaped dual-chain structure is formed by the oligomeric nucleic acid sequence and the one matched with it. Controlling the electric potential of the gene chip can further optimize hybridisation conditions. Its advantages are repeated application, low cost and the power for recognizing full match and single-point mismatch. The present sequence represents the gene chip associated oligonucleotide #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gene chip for recognizing full match and single-point mismatch, conhairpin-shaped probes having a detecting region and a stem region.
                                                                                                                                AAQ66708 standard; DNA;
                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 4 A; 3 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 6 (Disclosure); 11pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gene chip; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gene chip associated oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACD26289 standard; DNA; 15
            HHV6; Human Herpes Virus 6; Primers; Probes; polymerase chain reaction; ss.
                                                      Primer
                                                                                22-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-MAR-2002; 2002CN-00116302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-MAR-2002; 2002CN-00116302
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                                                                                                                                                                                                                            16
                                                                                                                                                                                                  4
                                                                                                                                                                                                                                                     12;
                                                   to amplify HHV6 derived sequences
                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wei F,
                                                                                                                                                                                                  ATGAACCGGAGG 15
                                                                                                                                                                                                                            ATGAACCGGAGG 27
                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                               (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hairpin; full match; single-point mismatch.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sun
                                                                             entry)
                                                                                                                                                                                                                                                                2.0%;
                                                                                                                                 16
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                                                                                                                                                                                                                                                                Score 12;
Pred. No.
                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                DB 7; Le
1.2e+05;
                        PCR; amplify;
                                                                                                                                                                                                                                                                             Length 15;
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RESULT 111
ACA58181/c
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Best Local &
                                                                                                                                                      29-MAR-1996;
20-OCT-1997;
19-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The inventors provide human Herpes useful for identification of HHV-6 are used in the invention
                                                                                                                                                                                                                                                                               Human; genotype determination; familial bipolar affective disorder; chromosomal region linked; locus associated with resistance; D4S402 D4S424; D4S431; D4S404; D11S394; D11S29; chromosome marker; primer;
                                                                                                                                                                                                                                                                                                                       Human familial bipolar affective disorder chromsome marker
                                                                                                                                                                                                                                                                                                                                             09-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                  ACA58181;
                                                                                                                                                                                                                                                                                                                                                                                       ACA58181 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 16 BP; 5 A; 2 C; 3 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1994-196175/24.
                         Determining a genotype associated with increased or decreased resistance to familial bipolar affective disorder in a family comprises determining the genotype of e.g., chromosomal regions D4S402 and D4S424.
                                                                                                                                                                                                                                        US2002192655-A1
                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HHV-6 derived nucleotide(s) - useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-MAY-1994
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        Disclosure;
                                                                      WPI; 2003-352708/33.
                                                                                                                                                                                               13-JUN-2001; 2001US-00881012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                            (GINN/) GINNS E I.
(EGEL/) EGELAND J A.
(PAUL/) PAUL S M.
                                                                                                                                                                                                                     19-DEC-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (KOKU-) KOKUSAI SHIYAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                              496 CATGAAAATTCA 507
                                                                                          ם,
                                                                                                                                                                                                                                                                                                                                                                                                                                          12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                         Egeland JA,
       Page 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                      96US-0014334P.
97US-0062924P.
98US-00175158.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.0%;
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                                                                                          Paul
        English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus
DNA. /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for identification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 6 derived nucleotide sequences AAQ66705-12 are primer set 1 (I),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 112
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Best Local
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                           The sequence is that of the hybridising region of tailed probe DRB83 for use in a method for determining HIA-DR beta sub-type in a nucleic acid sample. The method allows specific nucleic acid sequences of the second exon of HIA-DR beta genes to be amplified then probed for identification of polymorphic sequences. The amplified DNA is useful for typing the property of the amplified DNA is useful for typing the property of the amplified DNA is useful for typing the property of the amplified DNA is useful for typing the property of the property of the amplified DNA is useful for typing the property of 
                                                                                                                                                                                                                                                                                                                                                                                                                                    Method for determining HLA-DR beta sub-type in DNA sample - amplification and hybridisation with probes and primers, use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Erlich HA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
04-JAN-1993
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of polymorphic sequences. The ampl
homozgous or heterozygous samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-234644/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HLA-DR beta
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                                                                                                                                                                                                                                                                                                          Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Apple RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-DEC-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tissue typing; identity determination; disease susceptible;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sub-type tailed probe DRB83 hybridising region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Begovich
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(first entry)
                                                                                                                                                                                                                                                                                                          39; 90pp;
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                                                                                                                                                                                                                                                                                                          English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 T; 0 U; 0 Other;
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from a variety of sources
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Griffith RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .2e+05;
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BP; 3 A;

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RESULT 113
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Best Local
                                  The sequence is that of the hybridising region of tailed probe DRB61 for use in a method for determining HLA-DR beta sub-type in a nucleic acid sample. The method allows specific nucleic acid sequences of the second exon of HLA-DR beta genes to be amplified then probed for identification of polymorphic sequences. The amplified DNA is useful for typing homozogous or heterozygous samples from a variety of sources and for detecting allelic variants not distinguishable by serological methods. The typing system can be used in a reverse dot blot format which is simple and rapid to perform, produces detectable signals in minutes and can be utilised in tissue typing, determination of individual identity and identifying disease susceptible individuals. See also AAQ26092-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The typing system can be used in a reverse dot blot format which is simple and rapid to perform, produces detectable signals in minutes can be utilised in tissue typing, determination of individual identiand identifying disease susceptible individuals. See also AAQ26092-
                                                                                                                                                                                                                                                            Method for determining HLA-DR beta sub-type in DNA sample - amplification and hybridisation with probes and primers, use
                                                                                                                                                                                                                                                                                                                                      Erlich HA,
Apple RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                          06-DEC-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue typing;
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04-JAN-1993
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12; Conserv
                           (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             identity determination;
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Pred. No.
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1.2e+05;
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RESULT 114
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                                                                                                                                                                                                                                                                                                     The sequences given in AAQ92112-211 are probes which were used in the detection of a mutant p53 gene sequence. The DNA to be detected is amplified using PCR and then these probes which are pref. labeled using 32-P gamma-ATP are used to detect the mutant sequences. The primers and probes given in AAQ92098-219 are used in the method of the invention for detecting mammalian target DNA in sputum samples. Analysis of the target DNA is used to diagnose benign or malignant neoplasms of the lung. It is also useful for screening people at high risk or for monitoring progress of treatment of lung neoplasms. The method is based on the discovery that mutant target DNA associated with lung cancer is present at detectable levels in sputum. Cells shed into sputum from head and neck cancers may also be detected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting target nucleic acid in mammalian sputum - particularly for diagnosis of lung neoplasia involving mutation(s) in the K-ras oncogo or p53 tumour suppressor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primer; polymerase chain reaction; amplify; mutant; K-ras; PCR; flanking region; amplification; probe; detection; sputum; diagnosis; benign; malignant; neoplasm; lung; lung cancer; head; neck; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ92156 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               p53 detection probe, (codon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ92156;
Canine IL-2 receptor g-chain substrate position 1021
                               24-FEB-1999
                                                                                     AAV95212 standard; RNA; 17
                                                                                                                                                                                                                                                                            Sequence 17 BP; 4 A; 4 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 32; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-MAY-1995.
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                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1995-194114/25
                                                                                                                                                                                        490
                                                                                                                                                                                                                    12;
                                                                                                                                                           ທ
                                                                                                                                                             GGGCTGCATGAA 16
                                                                                                                                                                                        GGGCTGCATGAA 501
                                                                                                                                                                                                                   ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                             (first entry)
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                                                                                      ВP
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                                                                                                                                                                                                                                  Score 12;
Pred. No.
                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                  1.2e+05;
                                                                                                                                                                                                                                                 DB 2;
                                                                                                                                                                                                                                                 Length 17;
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                                                                                                                                                                                                                    Gaps
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AAV93889 to AAV94574 represent specifically claimed ribozymes, and AAV94575 to AAV95260 represent specifically claimed substrate sequences from the present invention. The ribozymes can be used for the treatment of, e.g. graft rejection, autoimmune disease, cancer, psoriasis, allergy and other inflammatory conditions. The ribozymes are also used to induce tolerance in a recipient to alloantigen from a donor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; IL-2 receptor g-chain; interleukin 2 receptor gamma chain; hammerhead ribozyme; hairpin ribozyme; substrate; expression; can autoimmune disease; psoriasis; allergy; inflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ribozymes targetted to interleukin autoimmune disease and allergies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      graft rejection; ss.
Sequence 17 BP; 3 A; 3 C; 10 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9824913-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Canis
                                                                                                                                                                                                                                                                           The present sequence invention describes ribozymes targeted to modulate the synthesis and/or expression of interleukin (IL)-2R gamma encoded RNA.
                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Page 47; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ъ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.
19
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Query Match
Best Local Similarity
Matches 12; Conserve
248 CCTGGAGCCCCT 259
                 Conservative
                         100.0%;
                 0
                          Score 12;
Pred. No.
                  Mismatches
                          1.2e+05;
                  0
                  Indels
                  0
                  Gaps
                  0
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2.0%;

DB 2;

Length 17

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맑
AAA18564 standard; RNA; 17
                                                                             13 CCTGGAGCCCCT 2
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Human TIE-2 substrate sequence SEQ ID NO:1790.

19-JUN-2000 AAA18564;

(first entry)

RESULT 116
AAA1854/c
ID AAA185
XX AAA185
XX AAA185
XX AAA185
XX Human
DT 19-JUN
DT 19-Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiotibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; ss. xippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss.

Homo sapiens

WO9950403-A2

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC hydrocarbon nuclear transporter (ARMT) gene, an integrin subunit beta 3 (C gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARMT, and AAA17168 to AAA1818 to AAA17168 to AAA1818 and AAA1918 to AAA1818 represent their corresponding target sequences; AAA18123 to AAA18123 to AAA1818 represent their corresponding target sequences; AAA181823 to AAA1818 represent their corresponding target sequences; AAA21689 to AAA21689 represent their corresponding target sequences; CC AAA21689 to AAA2188 represent their corresponding target sequences; CC AAA21689 to AAA2183 and AAA2276 to AAA2382 represent their corresponding target sequences. The ribozyme sequence for integrin subunit beta 3, and AAA2276 to AAA2382, AAA23343 to AAA2342 represent their corresponding target sequences. The ribozyme of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, CC integrin subunit beta 3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related consequences of tuberous sclerous, pot-wine stains, Sturge Weber and other syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, integrin subunit alpha-6, or integrin subunit beta-3, integrin subunit beta-3, integrin subunit beta-3, int
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 12
                                                                                                          Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinomathy, arthritic
                                                           dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis, age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration, psoriasis; verruca vulgaris; angiofibroma;
                                                                                                                                                                                                                                                                                                             19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                        AAA20707 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
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                                              tuberous
                                                                                                                                                                                                                                                         Integrin alpha 6 subunit substrate sequence SEQ ID NO:3933.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 integrin subunit alpha-6, or integrin subunit beta-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ribozymes for mRNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 CTGGAGCCCCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                   sclerosis; pot-wine stain; Sturge Weber syndrome
renaunay-Weber syndrome; Osler-Weber-Rendu syndron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGGAGCCCCTG 260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 4 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modulating the an angiogenic f
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Pred. No. 1.2e+0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                e synthesis, factors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 17;
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RESULT 118
AAV91124/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC hydrocarbon nuclear transporter (ARNIT) gene, an integrin submit beta 3 (2 gene, an integrin alpha 6 submit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNIT, and AAA17168 to AAA181955 to AAA19155 to AAA19155 to AAA19122 represent their corresponding target sequences; and AAA19155 to AAA19122 represent their corresponding target sequences; classified and AAA19155 to AAA1922 represent their corresponding target sequences; classified and AAA19155 to AAA1922 represent their corresponding target sequences; classified and AAA21689 to AAA21689 and AAA21689 to AAA21689 to AAA21689 to AAA21689 to AAA21689 to AAA21689 to AAA22336 to AAA22332 represent ribozyme sequence for integrin submit beta 3, and AAA22346 to AAA23362. AAA23342 represent their corresponding target sequences of the invention are used for modulating target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, classified to the invention are used for modulating the synthesis, especially ARNT, classified to the synthesis of the synthesis as well as capsicitions of tuberous scherosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, integrin submit alpha-6 or integrin submit alpha-6 or integrin submit alpha-6 or ARNT, Tie-2, integrin submit alpha-6 or integrin submit alpha-6 or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 7
           Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer inflammation; psoriasis; non-hepatic ascites; infection; genetic drift;
                                                                                                                           Human C-raf target site nucleotide position 1289
                                                                                                                                                                                                                    AAV91124;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17
                                                                                                                                                                         18-FEB-1999
                                                                                                                                                                                                                                                              AAV91124 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes enzymatic nucleic acid molecules with RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 55; Page 162; 305pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA encoding an angiogenic f
                                                                                                                                                                                                                                                                                                                                                                                                                       ω
5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               activity, which specifically cleave RNA encoded by
                                                                                                                                                                                                                                                                                                                                                                         UUUACCAAUUCA 17
                                                                                                                                                                                                                                                                                                                                                                                                                       TTTACCAATTCA 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                subunit alpha-6, or integrin subunit beta-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 A; 4 C; 0
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                                                                                                                                                                                                                                                              ΒP
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`•
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G; 0 T; 8 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12; DB 2;
Pred. No. 1.2e+05;
5; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Coeshott C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               synthesis, expression and/or stability
factors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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RESULT 119
AAA36309
ID AAA363
XX
AC AAA363
XX
DT 26-JUL
XX
DE Human
                                                                                                                                                                                                                                            S
                                                                                                                                                                                                                                                              CC capable of modulating a process in a biological system. The method CC comprises: (a) introducing into the system a random library of nucleic CC acid catalysts (NAC) having a substrate binding domain (SBD), comprising CC a random sequence, and a catalytic domain (CD); and (b) identifying NAC CC in systems where modulation has occurred and/or determining the sequence CC of at least part of the SBDs in such systems. Nucleic acid molecules with CC endonuclease activity and catalytic activity, from the present invention, CC are used to modulate gene expression in plant and mammalian cells and to CC cleave target nucleic acid, particularly for treating systemic diseases CC caused by specific RNA, e.g. cancer, inflammation, pooriasis, non-hepatic CC mutations in diseased cells and to determine c-raf RNA. Specifically NACs CC with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restences, psoriasis or rheumatoid arthritis, or CC generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and activity. AAV90922 to AAV93877 represent NACs that can be used in the method.
                                                                                                                                        멼
                                                                                                                                                                                                    Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A method has been developed for the identification of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 177; Page 149; 259pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-OCT-1997;
02-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JUL-1997;
22-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-MAY-1998;
                        26-JUL-2000
                                                                          AAA36309
                                                                                                                                                                                                                                          Sequence 17 BP; 5 A; 5 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-009494/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                 170
                                                                                                                                        12
                                                                                                                                                                                        l Similarity
12; Conserv
                                                                          standard; DNA; 17
                                                                                                                                        TGGAATTGCTCT 1
                                                                                                                                                                 TGGAATTGCTCT 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Beigelman L, N
J, Workman CT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matulic-Adamic J, Reynord:
Beigelman L, Mcswiggen JA,
I. Workman CT, Beaudry A,
                                                                                                                                                                                          Conservative
                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97US-0061321P.
97US-0061324P.
97US-0064866P.
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97US-0051718P.
97US-0056808P.
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                                                                                                                                                                                                    2.0%;
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                                                                                                                                                                                                    Score 12; DB 2;
Pred. No. 1.2e+
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                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Karpeisky A,
Sweedler D;
                                                                                                                                                                                                    1.2e+05;
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                                                                                                                                                                                                                Length 17
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                                                                                                                                                                                          Gaps
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Human genomic SNP allele specific oligonucleotide SEQ ID NO:375

06-JAN-2000

26-MAY-1999;

99WO-US011516

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RESULT 120
AAZ88059
ID AAZ880
XX AAZ880
XC AAZ880
XC AAZ880
DT 06-AUG
DT 20-APR
XX Lentiv
XX Lentiv
XX Gene e
XX Gen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     method comprises preparing a reduced complexity genomic sample. The genomic sample and analysing the RCG for the presence or absence of a SNP allele. The method can be used to characterise a tumour, to generate a genomic pattern for an individual genome or to generate a genomic code for a genome. The method can be used to assess whether a subject is at risk for developing a disease or to identify a set of SNP alleles associated with a disease. The method can also be used to perform linkage analysis. AAA35944 to AAA35947 represent sequences used in the exemplification of the present invention. AAA35948 to AAA36632 represent nucleotide seminars.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; single nucleotide polymorphism; SNP; genotyping; DNA analysis; allele specific oligonucleotide; ASO; redece complexity genome; RCG; genomic classification; identification; DNA fingerprinting; tumour characterisation; hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A method has been developed for detecting the presence or absence single nucleotide polymorphism (SNP) allele in a genomic sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detection of single nucleotide polymorphisms in genomes by preparation and analysis of reduced complexity genomes, useful for genotyping, fingerprinting and determining allele frequency of SNPs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 4 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 64; 111pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-293181/25.
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                                                                                                                                                                                                                                                                                                                                                         Lentiviral vector; packaging; gag; pol; gene therapy; infection;
gene expression; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lentiviral proviral long terminal repeat DNA PCR primer #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-AUG-2003
20-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ88059;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Landers JE,
                                                                                                                                                                                   WO200000600-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MASI ) MASSACHUSETTS INST TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                409
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12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCCATCATGACC 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0101757P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 3; Lt
1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>.</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of a
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AAZ88115
ID AAZ8
XX
AC AAZ8
AC AAZ8
AX Lent
XX Lent
XX Gen
XX Synt
XX WO2/
XX WO2/
XX WO2/
XX E Cha
PD 06-,
XX 26-|
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              produce packaging vector particles comprising functional Gag and Pol proteins. The vectors are useful for transforming (eukaryctic) cells to express specific genes at high levels, e.g. for gene therapy. The improved vectors are safer, yet permit increased efficiency of packaging the recombinant viral genome and increased long-term gene expression. These properties are required for gene therapy as a means of treating infectious and non-infectious diseases. Unlike other retroviruses, the lentiviruses are able to infect non-dividing cells. The present sequence represents a lentiviral proviral long terminal repeat DNA PCR primer which is used in the exemplification of the present invention. (Updated on 06-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleotide sequence encoding Gag and Pol proteins of a reference lentivirus that differs from the reference lentivirus at least in that:

(a) its major splice donor site is either deleted or is sufficiently different from the reference lentivirus so that it is not a potential site for homologous recombination; and (b) it lacks a functional major packaging signal so that the introduced vector causes the host cell to
                                                                                                                                                                                                                                                                                                                                                                                               Lentiviral vector; packaging; gag; pol; gene therapy; infection; gene expression; PCR primer; ss.
  WPI; 2000-137067/12
                                                                                                                                                                                                                                                                                                   WO200000600-A2
                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hirt DNA PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New packaging vector comprising a nucleotide sequence encoding Gag and Pol proteins of a reference lentivirus useful for the delivery of non-lentiviral genes to target cells.
                                                   Chang L;
                                                                                                                                                    26-MAY-1998;
                                                                                                                                                                                                  26-MAY-1999;
                                                                                                                                                                                                                                                  06-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ88115 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a packaging vector (PV) comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 132; 311pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chang L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-MAY-1998;
                                                                                                   (CHAN/) CHANG
                                                                                                                                                                                                                                                                                                                                                                                                                        mentiviral vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CHAN/) CHANG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTCCTGGAGCCC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 3 A; 7 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                    98US-0086635P
                                                                                                                                                                                                  99WO-US011516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98US-0086635P.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 3; Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
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New packaging vector comprising a nucleotide sequence encoding Gag and PT Pol proteins of a reference lentivirus useful for the delivery of non-PT lentiviral genes to target cells.

XX

Example 200; Page 213; 311pp; English.

XX

The present invention describes a packaging vector (PV) comprising a CC nucleotide sequence encoding Gag and Pol proteins of a reference CC lentivirus that differs from the reference lentivirus at least in that:

CC (a) its major splice donor site is either deleted or is sufficiently CC different from the reference lentivirus so that it is not a potential concentration of the reference lentivirus so that it is not a potential concentration of a reference concentration of a reference concentration of the proteins. The vector recombination, and (b) it lacks a functional major concentration of the recombinant viral genome at high levels, comprising functional Gag and Pol concentrations. The vectors are useful for transforming (eukaryotic) cells to express specific genes at high levels, e.g. for gene therapy. The comprises specific genes at high levels, e.g. for gene therapy. The compress specific genes and non-infectious diseases. Unlike other means of treating confectious and non-infectious diseases. Unlike other retroviruses, the lentiviruses are able to infect non-dividing cells. The present sequence represent sequence paresent invention

XX

Sequence 17 BP; 3 A; 7 C; 5 G; 2 T; 0 U; 0 Other;

Query Match

2.0%; Score 12; DB 3; Length 17;
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Query Match 2.0%; Score 12; DB 3; Length 17;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

246 CTCCTGGAGCCC 257
||||||||||||||
5 CTCCTGGAGCCC 16
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RESULT 122
AAZ88064
ID AAZ88064 standard; DNA; 17 BP.

XX
AC AAZ88064;
XX
DT 20-APR-2000 (first entry)
XX
DE MLV proviral long terminal repeat DNA PCR primer #2.

XX
Lentiviral vector; packaging; gag; pol; gene therapy; infection;
XX
KW gene expression; PCR primer; ss.

XX
VS
Murine leukemia virus.

XX
VX
VO00000600-A2.
XX
VG-JAN-2000.
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New packaging vector comprising a nucleotide sequence encoding pol proteins of a reference lentivirus useful for the delivery

Gag and

WPI; 2000-137067/12.

Chang L;

26-MAY-1999; 26-MAY-1998;

99WO-US011516 98US-0086635P

(CHAN/) CHANG

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The present invention describes a packaging vector (PV) comprising nucleotide sequence encoding Gag and Pol proteins of a reference

Example; Page 132; 311pp; English.

entiviral genes to target cells.

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RESULT 123
ABK02421/c
Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; humaning disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                      11-FEB-2000; 2000US-0181797P
28-FEB-2000; 2000US-0185516P
06-MAR-2000; 2000US-0187128P
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(a) its major splice donor site is either deleted or is sufficiently
different from the reference lentivirus so that it is not a potential
                                      WPI; 2001-607195/69
                                                                                                                                       (RIBO-)
(BLAT/)
(MCSW/)
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                                                                                                                                                                               RIBOZYME PHARM INC
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CHOWRIRA B M
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Pred. No.
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constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and
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Claim 88; Page 132; 200pp; English

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down considerably most acids acids may be enzymatic nucleic acids (e.g. a ribozyme or a composed acids may be enzymatic nucleic acids (e.g. a ribozyme or a composed and individual considerably motif) processessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) processessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) processessing an NCH motif). The CD20-targetting nucleic acid is used to cleaver RNA composed in the presence of a divalent cation that is preferably Mg^2+. CC Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more considerables. In particular, the CD20 targetting nucleic acid may be used to cleave RNA composed in the composed in the level of CD20. The treatment may further comprise the use of one or more considerable in the composed in the considerable in the composed considerable in the considerable in th 8 an amberzyme molecule of the invention

Sequence 17 BP; 1 A; 6 C; 6 G; 0 T; 4 U; 0 Other;

S Matches Query Match Local з62 стемесссемее 373 1 Similarity 12; Conserv Conservative 100.0%; 2.0%; .. Score 12; DB 4; L Pred. No. 1.2e+05; Mismatches 0 Length 17; 0 Gaps 0

ABK00051 standard; RNA; 17 12-MAR-2002 ABK00051; (first entry) Въ

Human NOGO Hammerhead Ribozyme #51.

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CTGAGCCCGAGG 4

RESULT 124
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ID ABKO00
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Homo s Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NGG0; hammerhead ribozyme; NMzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; b-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; NCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkitsen's disease; ataxia; Huntingcon's disease; Creutzfeldt-Jakob disease; muscular neurodegenerative disease

acid molecules,

e.g., enzymatic nucleic acids

and

Synthetic

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down CC regulates expression of a neurite growth inhibitor gene (NOGO). The CC nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CD DNazyme) an Inazyme (an endolytic nucleic acid cleaving an RNA molecule CD possessing an NCH motif), a C-cleaver (cleaving RNA with a NYN motif) processing an molecule contact of the collaryme (cleaving RNA with an NGH triplet), a zinzyme (cleaving RNA cOC with a YGY motif). The CD20-targetting nucleic acid is used to cleaver RNA CC with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA CC with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA CC with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA CC the cell and treat a patient having a condition associated with the level of CD20 in the treatment may further comprise the use of one or more contacted with a cell to reduce CD20 activity of the cell and treat a patient may further comprise the use of one or more contacted with in the level of immune thrombocytopaenia, and inflammatory arthropathy. The NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO activity of the nucleic acid may be contacted with a cell to reduce NOGO activity of the nucleic acid may be contacted with a cell to reduce NOGO activity of the nucleic acid may be used to cleave RNA of the NOGO activity of the nucleic acid may be used to cleave the use of one or more cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more cell and treat a patient having a condition associated with the level of CNGO. The treatment may further comprise the use of one or more cell and treat a patient having a condition associated with the level of CNGO. The NOGO activity of the condition accidence of a collar, the NOGO activity of the nucleic acid may be used to cleave the NOGO activ
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Best Local :
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28-FEB-2000; 2000US-0185516P
06-MAR-2000; 2000US-0187128P
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                                                                                                                                                          BP; 1
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Pred. No. 1.2e+05;
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RESULT 125 ABK00947/c

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The invention relates to a nucleic acid molecule which down regulates CC expression of a CD20 gene and a nucleic acid molecule which down CC regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids (e.g. a ribozyme or a nucleic acids (e.g. a ribozyme or a CC DNAzyme) an Inozyme (an endolytic nucleic acids cleaving an RNA molecule cossessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) preceded in the YGY motif). The CD20-targetting nucleic acid is used to cleave RNA cC with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA cC with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA cC with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA cC the cell and treat a patient having a condition associated with the level cof CD20. The treatment may further comprise the use of one or more comparated to the cell and treat a patient having a condition associated with the level composition of the lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-cC Hodgkin's lymphoma (NHL), bulky low-grade or follicular NHL, lymphocytic lymphoma (NHL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, cell to reduce CP (lymphoma, leukaemia, and inflammatory arthropathy. The NOGO-compresence of a divelent cation that is preferably MG'2'+. Furthermore, the concleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of collar not treatment may further comprise the use of one or more
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28-FEB-2000; 2000US-0185516F
06-MAR-2000; 2000US-0187128F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapies. In particular, the NOGO-targetting nucleic acid may be used treat central nervous system (CNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), Parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present
                                                                         Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurit growth inhibitor gene useful for treating, e.g., lymphoma, leukemia
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28-FEB-2000; 2000US-0185516P
06-MAR-2000; 2000US-0187128P
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. 1.2e+05;
                                                                          a CD20 gene or neurite
g., lymphoma, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                            and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ť
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Claim 88; Page 132; 200pp; English

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nucleic acid molecule which down regulates

09-FEB-2001; 2001WO-US004273

WO200159103-A2

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cregulates expression of a CD20 gene and a nucleic acid molecule which down cregulates expression of a neurite growth inhibitor gene (NOGO). The CD NAZyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a consistency and inozyme (an endolytic nucleic acid cleaving an Index motif) processing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) processing an NCH motif). The CD20-targetting nucleic acid is used to cleaver RNA consistency, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-cell two processes and the cell lymphoma (NHI), bulky low-grade or follicular NHI, lymphocytic lymphoma (MCI), immunocytoma (IMC), small B-cell lymphocytic lymphoma (MCI), immunocytoma (IMC), small B-cell lymphoma, the NGO-cell lymphoma (MCI), immunocytoma (IMC), small B-cell lymphocytic lymphoma (MCI), immunocytoma that is preferably MC 2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NGO gene in the cell and treat a patient having a condition associated with the level of the ragetting nucleic acid may further comprise the use of one or more therapies. In particular, the NGO-targetting nucleic acid may be used to cleave RNA of the NGO activity of the cell and treat a patient having a condition associated with the level of the cell and treat a patient having a condition associated with the level of the cell and treat a patient having a condition associated with the level of the cell and treat a patient having a condition associated with the level of cell and treat a patient having a condition associated with the level of the cell and treat a patient action of NGO expression. The present cell and treat which respond to the modulation of NGO expression. The present sequence is an amberzyme molecule of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                          Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NCG0; hammerhead ribozyme, DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin; slymphoma; NHI; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHI; mantle-cell lymphoma; NCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkingon's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human NOGO Zinzyme #168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK01846;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK01846 standard; RNA; 17
                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                              Creutzfeldt-Jakob disease; muscular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    362 СТСАССССАСС 373
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTGAGCCCGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 1 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 4; Le . 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                                                           neurodegenerative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                     ribozyme;
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RESULT 128
ABA77345
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC DNAzyme) and incoryme (an endolytic nucleic acid cleaving an NYM molecule CC possessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) preceded an amberzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA CC with a YGY motif) The CD20-targetting nucleic acid is used to cleave RNA CC with a YGY motif) The CD20-targetting nucleic acid is used to cleave RNA CC with a YGY motif) that is preferably Mg^2^+. CF Purthermore, it may be contacted with a cell to reduce CD20 activity of CC cleaver. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular NHL, lymphore (CC treat lymphoma (NHL), bulky low-grade or follicular NHL, lymphore, leukaemia, B-cell lymphoma, low-grade or follicular NHL, lymphore, cell conditions (NGC), immune thrombocytopaenia, and inflammatory arthropathy. The NGCO gene in the CC targetting nucleic acid may be contacted with a cell to reduce NGCO activity of the CC uncleic acid may be contacted with a cell to reduce NGCO activity of the CC uncleic acid may be contacted with a cell to reduce NGCO activity of the CC uncleic acid may be contacted with a cell to reduce NGCO activity of the CC uncleic acid may be contacted with a cell to reduce NGCO activity of the CC uncleic acid may be used to cleave RNA of the NGCO activity of the CC uncleic acid may be used to creat price of a disease, and inflammatory arthropathy. The NGCO activity of the CC uncleic acid may be contacted with a cell to reduce NGCO activity of the CC uncleic acid may be used to cleave RNA of the NGCO activity of the CC uncleic acid may be used to creat central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), CC chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), Carkinson's disease, ataxia, Huntington's disease, Creutzfeld-Jakob CC disease, which respond to the modulation of NGCO expression. T
                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-FEB-2000; 2000US-0181797P.
28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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(BLAT/)
(MCSW/)
p53 mutation correcting oligonucleotide SEQ ID NO: 191
                                             24-JAN-2002
                                                                                                                                  ABA77345 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 88; Page 98; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  central nervous system injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-607195/69.
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                                                                                                                                                                                                                                                                                           362
                                                                                                                                                                                                                                                14 CTGAGCCCGAGG 3
                                                                                                                                                                                                                                                                                                                                  12;
                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RIBOZYME PHARM INC
BLATT L.
MCSWIGGEN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CHOWRIRA B M.
                                                                                                                                                                                                                                                                                      CTGAGCCCGAGG 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      is a zinzyme molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                    Conservative
                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                      2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                      6 C; 6 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                  Score 12; DB; Pred. No. 1.2; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                      DB 4; Le
, 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                  RESULT 129
ABA77317
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ABA77317; 24-JAN-2002

(first entry)

ABA77317 standard; DNA; 17

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ATGAACCGGAGG

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                                                                                                                              The present invention provides single-stranded oligonucleotides which can CC be used for the targeted alteration of genomic sequences, where the CC oligonucleotide has at least one mismatch compared with the genomic CC sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, cretinoblastoma, BRCA1, BRCA2, CFFR, cyclin-dependent kinase inhibitor 2A CC (CDKN2A), APC, Pactor V, Factor VIII, Factor IX, haemoglobin alpha locus 2 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, applipoprotein E (APCE), LDL receptor (LDLR), UDP-glucuronosyltransferase CC (UGTI), amyloid precursor protein CC (UDRA), UPP-glucuronosyltransferase CC (UGTI), amyloid precursor protein be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, CC haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, CC haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, CC Alzheimer's disease. The present sequence is one of the gene correcting CC oligonucleotides of the invention
                                           Matches
                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UCFT; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzhimer; disease; cytostatic; antisickling; antianaemic; haemostatic; Alzhimer; disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                         Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-MAR-2000; 2000US-0192176P.
27-MAR-2000; 2000US-0192179P.
01-JUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 7; Page 54; 294pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kmiec EB, Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-MAR-2001; 2001WO-US009761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antilipemic; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYDE ) UNIV DELAWARE
                                                               Local
  16 ATGAACCGGAGG 27
                                         l Similarity
12; Conserv
                                           Conservative
                                                                                                                         BP; 5 A; 6 C; 5 G; 1 T; 0 U; 0 Other;
                                                               100.0%;
                                                                                    2.0%;
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                                           0
                                                               Score 12;
Pred. No.
                                           Mismatches
                                                                                  DB 4;
                                                                 1.2e+05;
                                                                                  Length 17;
                                           Indels
                                           0
                                           Gaps
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p53 mutation correcting oligonucleotide SEQ ID NO: 163

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RESULT 130
ABA77318/c
ID ABA773
XX
AC ABA773
XX
AC ABA773
XX
AC ABA773
XX
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                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                     the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, Cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein E (APCB), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGT1), amyloid precursor protein (APC), presentiin-1 (PSEN1) and presentiin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCAL; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2h; ChNU2h; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilla; alpha thalassaemia; haemoglobin alpha locus 1; MLHI; APOE; mismatch repair; MSH4; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UCT1; syndrome; APP; PSENI; antisense; UDP-glucuromosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention provides single-stranded oligonucleotides which obe used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-MAR-2000; 2000US-0192176F.
27-MAR-2000; 2000US-0192179F.
01-JUN-2000; 2000US-0208538F.
30-OCT-2000; 2000US-0244989F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide for targeted alterations of genetic sequences treating cystic fibrosis, comprises at least one mismatch and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200173002-A2
                                                                                                                                                                                                                                                                                                     Sequence 17
                                                                                                                                                                                                                                                                                                                                         oligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-639230/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYDE ) UNIV DELAWARE
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                                                                       ABA77318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7; Page 51; 294pp; English
                                                                                                                                                                                               16
                                                                                                                                                                                               ATGAACCGGAGG
                                                                       standard;
                                                                                                                                                              ATGAACCGGAGG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gamper HB,
                                                                                                                                                                                                                                                                                                      BP; 4 A;
                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                 2.0%;
                                                                       DNA;
                                                                                                                                                                                                                                                                                                      6 C; 6 G; 1 T; 0 U; 0 Other;
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                                                                         17
                                                                                                                                                                                                                                Score 12; DB; Pred. No. 1.2
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N
                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                 DB 4; L
                                                                                                                                                                                                                                                                  Length 17;
                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and for
chemical
                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               can
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Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR, cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltramsferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mutation correcting oligonucleotide SEQ ID NO: 164.
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antilipemic; ss.

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27-MAR-2000; 2000US-0192176P.
27-MAR-2000; 2000US-0192179P.
01-JUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
                                                                                                                                                                                                                                                 04-OCT-2001
                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                    WO200173002-A2
                                     Kmiec EB,
                                                                                                                                                                                                           27-MAR-2001; 2001WO-US009761.
                                                                         (UYDE ) UNIV
2001-639230/73
                                   Gamper HB,
                                                                           DELAWARE
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Oligonucleotide modification. Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemica chemical

Claim 7; Page 52; 294pp; English

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the coligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus (CHBA1), haemoglobin alpha locus 2 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH, MSH2, WSH6, amploid precursor protein (ADC), Dresenilin-1 (FSEN) and CC (UGT1), amyloid precursor protein (APC), presenilin-1 (FSEN) and CC presenilin-2 (FSENY2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, CC haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, CC harbonier's disease, melanoma, adenomatous polyposis of the colon and CC various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention

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Sequence 17
BP; 1 A;
 6 C; 6
 G; 4 T; 0 U; 0 Other;
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            Ś
                           Matches
                                        Query Match
                                 Local
15
             16 ATGAACCGGAGG
                          l Similarity
                          Conservative
                                 100.0%;
             27
                                        2.0%;
                           0;
                                 Score 12;
Pred. No.
                           Mismatches
                                 DB 4;
                                 .2e+05;
                                        Length 17;
                           0
                           Gaps
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0

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RESULT 131
ABA77346/c
ID ABA773
XX
AC ABA773
XX
           ABA77346
                                ABA77346 standard; DNA; 17 BP
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24-JAN-2002

(first

entry)

<u>р</u>53

correcting oligonucleotide SEQ ID NO:

24-JAN-2002

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                 RESULT 132
                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                         oligonuclectide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFFR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein E (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UCT1), amyloid precursor protein (APC), presentlin-1 (PSEM1) and presentlin-2 (PSEM2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemogbhilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UCT1; syndrome; APP, PSENI; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentlin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                          Sequence 17 BP; 1 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
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27-MAR-2000; 2000US-0192179P.
01-UUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
                                                                                                                                                                                                                                                               various syndromes. To oligonucleotides of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 7; Page 54; 294pp; English
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Sequence 17

BP; 7

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and/or amount specifically of hgDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hgDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hgDMLP-1 may be used for diagnosing a disorder associated with the expression of hgDMLP-1, in particular heart and skeletal muscle disorders. hgDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hgDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
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30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                          protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 provide initial substrates for the recombinant engineering of hGDMLP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                                                                       protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 7365; 214pp; English.
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27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                      invention describes a human genome-derived myosin-like
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                                                                                                       protein i (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration
therapeutic supplement in patients having specific deficiency in hGDMI production, and in vaccines or for replacement therapy. The production, sequences encoding hGDMLP-1 may be used for diagnosing disorder associated with the expression of hGDMLP-1, in particular heal and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.
                                                                                                                                                                                                                                                                                                                New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
                                                                            and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as
                                                                                                                                                                                                                                                                                   Disclosure;
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2001WO-US000667.
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disorder; amplicon; screening; ss.
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                                                                                                                                                                                                 New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                           Shannon
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP 1 can be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to

for the

recombinant

ę,

hGDMLP-1

hGDMLP-

Disclosure;

SEQ ID NO

7367; 214pp; English

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expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP C-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as characteristic capture probes for surface-enhanced laser desorption ionisation, as characteristic capture probes for surface-enhanced laser desorption ionisation, as characteristic capture probes for surface-enhanced laser desorption ionisation, as characteristic deficiency in hGDMLP-1 capture probes for surface encoding hgDMLP-1 may be used for diagnosing a construction, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hgDMLP-1 may be used for diagnosing a construction of hgDMLP-1, in particular heart can disorder associated with the expression of hgDMLP-1, in particular heart can skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the exemplification of the present invention. N.B. captures data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO as format/min/minished part of the present directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Matches 12
WPI; 2002-179446/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7369.
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                                                                                                                                                                                                                                                                                                                                                                       30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234687P.
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                                                                                                                         (AEOM-) AEOMICA INC.
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; 2001WO-US000664.
2001WO-US000666.
; 2001WO-US000666.
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CC provide initial substrates for the recombinant engineering of hGDMLP-1 couldeld acids in samples, as amplification substrates, to comprove the variants having desired phenotypic improvements, and for comprovered the proteins having desired phenotypic improvements, and for compressing the proteins. The hGDMLP-1 proteins or polypeptides may be compressed as immunogens to raise antibodies that specifically recognise hGDMLP conditions of the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 capture probes for surface-enhanced laser description ionisation, as considered to the replacement therapy. The plynucleotide sequences or for replacement therapy. The plynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a consider associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 may be used for diagnosing a considered sequence represents an oligomer used in the screening of the present sequence represents an oligomer used in the screening of the present sequence of the exemplification of the present invention. N.B.

The sequence data for this patent did not form part of the printed confidence of the printed confidence
Sequence 17 BP; 7 A; 1 C; 4 G; 5 T; 0 U; 0 Other;
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desorption
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              as specific biomolecule capture probes for surface-enhanced laser sorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                       ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              be used in gene therapy and vaccine production. The hgDMLP-in acids can be used as probes to defeat the hgrand production.
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Matches
                                            Query Match
Best Local 9
                 499 GAAAATTCAGTT 510
1 GAAAATTCAGTT 12
                                   12;
                                            Similarity
                                  ilarity 100.0%;
Conservative
                                                     2.0%;
                                           Score 12; ; Pred. No.
                                   0;
                                   Mismatches
                                  1.2e+05;
hes 0;
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RESULT 136
Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence
                                          29-MAY-2002
                                                                                 ABN07372
                                                                                                                     ABN07372 standard; DNA; 17
                                      (first entry)
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muscle; myosin; chromosome skeletal muscle disorder; a Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; e 22; gene amplicon;

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30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
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21-SEP-2000;
27-SEP-2000;
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                                                  2001WO-US000667
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New polypeptide,

for raising antibodies

that

hGDMLP-1 proteins,

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Query Match
Best Local Similarity
Thes 12; Conserve
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The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed and the sequence data for this patent did not form part of the printed and the sequence data for this patent did not form part of the printed and the sequence where the sequence is the patent of the printed and the sequence data for this patent did not form part of the printed and the sequence where the sequence contains the sequence that for this patent did not form part of the printed and the sequence where the sequence contains the sequence that section is patent of the printed and the sequence where the sequence contains the sequence contains and the sequence
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30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001US-0266860P.
                                                                                                                                                                                                                                                        Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; hear muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                                                                                                                                                                                Human GDMLP-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABN07374 standard; DNA; 17 BP
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26-MAY-2000; 2000US-0207456P
                                                25-MAY-2001; 2001WO-US016981.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                hGDMLP-1; heart;
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CC inspection, and in vaccine production. The inspirate of the protection acids can be used as probes to detect, characterise and quantify of hGDMLP-1 nucleic acids in samples, as amplification substrates, to comprove the initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for carpessing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP comprovements, and in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as concentration, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The groynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a complete disorders. hGDMLP-1 is localised to chromosome 22. The present sequence with the expression of hGDMLP-1, in particular heart cand skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the exemplification of the present invention. N.B. CC hGDMLP-1 sequence data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for this pattent did not form part of the printed complete data for the print
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-179446/23.
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27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rotein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-can be used in gene therapy and vaccine production. The hGDMLP-1
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2001WO-US000665.
2001WO-US000666.
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2001US-0266860P
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2001WO-US000669.
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2000US-0236359P.
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Sequence
 17
BP;
7 A;
 2 C; 4 G; 4 T; 0 U; 0 Other;
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Matches
               Query Match
         Local
l Similarity 100
12; Conservative
         100.0%;
                  2.0%;
Score 12; DB;
; Pred. No. 1.2;
0; Mismatches
0
         DB 6; Le
1.2e+05;
                 Length 17
 Indels
0
 Gaps
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RESULT 138
ABV85812
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ABV85812 standard; DNA; 17 BP
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499 GAAAATTCAGTT 510

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                    Human pp-GaNTase 10 scanning 17-mer SEQ ID NO:805.
                                    11-DEC-2002 (first entry)
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ABV85812

Human; UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase pp-GaNTase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; sca 10;

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RESULT 139
ABV85811
ID ABV858
XX
AC ABV858
XX
AC ABV858
XX
DT 11-DEC
XX
DE Human
XX
KW Human;
KW pp-GaN
XX
XX
SS.
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Matches 12
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30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
23-MAY-2001; 2001WO-US00064761.
30-AUG-2001; 2001US-0315984P.
                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an isolated nucleic acid (I) encoding a human UDP-GalNA:polypeptide N-acetylgalactosaminyltransferase 10 (pp-GaNTase 10, EC 2.4.1.41) protein. Human pp-GaNTase 10 is located to chromosome 7q11.2. (I) can be used in gene therapy, Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of pp-GaNTase. The sequence given in Abecreased expression and ABP53502 to ABP53504 are given in the exemplification of the present invention. N.B. The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid encoding human UDP-GalNAc:polypeptide N-cetylgalactosaminyltransferase 10 protein is useful to diagnose, prevent and treat disorders associated with reduced or over expression of the
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30-JAN-2001;
                                       pp-GaNTase
                                                        Human;
                                                                                                           11-DEC-2002
                                                                                                                                                               ABV85811 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; SEQ ID NO 805; 59pp; English.
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                                                                             pp-GaNTase 10 scanning 17-mer SEQ ID NO:804.
                                      UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10;
Tase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning;
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Pred. No.
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Homo sapiens

Synthetic.

Homo sapiens

EP1243660-A2

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RESULT 140
ABV85814
ID ABV858
XX ABV858
XX ABV858
XX I1-DEC
XX Human
XX Human
XX Human
XX EW Human
XX Spr-GaN
KW Sp-GaN
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XX Synthe
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30-JAN-2001;
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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                                                                             pp-GaNTase 10;
                                                                                                                    Human pp-GaNTase
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                                                                                                                                                                                                                                                                                                                                                                          Sequence 17
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                                                                                            Human;
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                                                                                                                                                                                                                                                                                                                       12;
                                                                          UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10;
Tase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning;
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2001WO-US000666.
2001WO-US000668.
2001WO-US000668.
2001WO-US000669.
2001WO-US000679.
2001US-00864761.
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
23-MAX-2001; 2001WO-US00064761.
30-AUG-2001; 2001US-0315984P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated nucleic acid (1) encoding a human UDP-GalNA:polypeptide N-acetylgalactosaminyltransferase 10 (pp-GaNTase 10, EC 2.4.1.41) protein. Human pp-GaNTase 10 is located to chromosome 7q11.2. (1) can be used in gene therapy. Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of pp-GaNTase. The sequence given in the decreased expression and ABP53502 to ABP53504 are given in the exemplification of the present invention. N.B. The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid encoding human UDP-GalNAc:polypeptide N-cetylgalactosaminyltransferase 10 protein is useful tand treat disorders associated with reduced or over e
                                          EP1243660-A2
                                                                               Homo sapiens
                                                                                                                     pp-GaNTase 10;
                                                                                                                                     Human;
                                                                                                                                                            Human pp-GaNTase 10 scanning 17-mer SEQ ID NO:808.
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Fase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning;
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RESULT 142
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid encoding human UDP-GalNAc:polypeptide N-cetylgalactosaminyltransferase 10 protein is useful to diagnose, prevent and treat disorders associated with reduced or over expression of the
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30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
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                                                                                                          Homo sapiens
Synthetic.
                                                                                                                                                                                                       Human pp-GaNTase
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                           25-JAN-2002; 2002EP-00001161
                                                                                 EP1243660-A2
                                                                                                                                                               Human; UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10; pp-GaNTase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanni
                                                                                                                                                                                                                                    11-DEC-2002
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23-MAY-2001;
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                                                      25-SEP-2002
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12; Conserv
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; 2001WO-US000669.
; 2001WO-US000670.
; 2001US-00864761.
; 2001US-0315984P.
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Pred. No. 1.2e+05;
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30-JAN-2001; 2001WO-US000663

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RESULT 143
ABV65813
ID ABV6581
XX ABV6580
XX ABV6580
XX II-DEC
XX Human
XX Human
XX Human
XX Epl-GaN
KW Ep.-GaN
KW Ep.-Ga
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Best Local S
Matches 12
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30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
23-MAY-2001; 2001WO-US00064761.
30-AUG-2001; 2001US-0315984P.
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  30-JAN-2001; 2001WO-US000663
30-JAN-2001; 2001WO-US000664
30-JAN-2001; 2001WO-US000666
30-JAN-2001; 2001WO-US000666
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                                                                                                                                                                                                                                                                          Synthetic
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Pred. No.
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1.2e+05;
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RESULT 144
ABV79548/c
ID ABV795
XX ABV795
XX ABV795
XX Human
XX Human
XX Human
XX Human
XX Homo s
XX For Don Or-Auc
XX EP1229
XX EP1229
XX EP1239

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30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
23-MAY-2001; 2001US-00864761.
30-AUG-2001; 2001US-0315984P.
                                                30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10 (pp-GaNTase 10, EC 2.4.1.41) protein. Human pp-GaNTase 10 is located to chromosome 7q11.2. (I) can be used in gene therapy. Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of pp-GaNTase. The sequences given in ABV85011 to ABV86689 and ABP53502 to ABP5304 are given in the exemplification of the present invention. N.B. The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied by the European Patent Office
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                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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12; Conserv
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2001WO-US000664.
2001WO-US000665.
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                                                                                                                                                                                                                                                                                                                                                                                                                          muscle; colon; male infertility;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>,</u>
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 6;
1.2e+(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .2e+05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to diagnose, prevent expression of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17
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30-JAN-2001; 23-MAY-2001;

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RRSULT 145
ABV79554/c
ID ABV795
XX ABV795
XX ABV795
XX
DE Human;
XX Human;
XX Human;
XX Human;
XX Prosta
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stopy codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL gene was important for diagnosing a discreter caused by mutation in HTPL, and in the thorapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of such discreter sinclude discreters of testis, or darenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and agents for the life was used in the second of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 167; 718pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-676582/73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human HTPL scanning oligonucleotide SEQ ID 800.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABV79554 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human testis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABV79554;
                                                                                                                                                                                                                             28-JAN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infertility and cancer. The present oligonucleotide was used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; testis expressed Patched like protein; testis; adrenal; liver; testis expressed Patchen emarrow; brain; kidney; lung; placenta; erm cell development; bone marrow; brain; kidney; lung; placenta; es; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           l Similarity
12; Conserv
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; 2001WO-US000663.
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; 2001WO-US000665.
; 2001WO-US000667.
; 2001WO-US000668.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 2 A;
                                                                                                                                                                                                                             2002EP-00001167.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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two. One of the single base pair differences between the ctwo. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL CC shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10pl2.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in cc therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human cf fortal liver, bone marrow, brain, kidney, lung, placenta, prostate, selectal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001US-00864761.
09-OCT-2001; 2001US-0327898P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and
Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-676582/73
                                              example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 168; 718pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or treating subjects having defects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (AEOM-) AEOMICA INC
  B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                              invention relates to human testis expressed Patched like
PL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). H
  2 A;
  4 C; 8
  G; 3 T; 0 U; 0 Other;
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멼
                                 Matches
                                         Query Match
Best Local
                                Local Similarity
mes 12; Conserv
               385
12
CIGCACCGCGCC 1
               CTGCACCGCGCC 396
                                 Conservative
                                        100.0%;
                                                2.0%;
                                Score 12; DB;
; Pred. No. 1.2;
0; Mismatches
                                 0
                                        DB 6; L
                                 <u>..</u>
                                                Length 17;
                                  Indels
                                 0
                                 Gaps
                                  0
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ACC53318
                                                                                                                       ss; tumour suppressor; antitumour; cytostatic; tumour suppression; tumour regression; apoptosis; virus resistance; diagnosis; cellular degeneration.
                                                                                                                                                               Human tumour
                                                                                                                                                                                  27-JUN-2003
                                                                                                                                                                                                                       ACC53318 standard; DNA; 17 BP
                           20-JUN-2001; 2001FR-00008139.
                                              20-JUN-2001; 2001FR-00008139
                                                                 27-DEC-2002
                                                                                   FR2826373-A1.
                                                                                                     Homo sapiens
                                                                                                                                                             suppressor sequence #2085.
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(MOLE-) MOLECULAR ENGINES LAB SA

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Best Local
given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence containing at least 15 consecutive alignment, at least 80 % identity to the 17 mer sequence, a sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for account.
                                                                                                                                                       New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABT37905 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           with tumour suppression or regression, apoptosis or virus resist invention relates to these sequences or sequences having at least identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. To invention is used to diagnose or treat viral disease or disease characterized by development of tumour cells or cellular degener
                                                                                           The invention relates to a novel isolated 17 mer nucleic acid sequence
                                                                                                                           Disclosure; Page 448; 720pp; French.
                                                                                                                                                                                                                                                                                                           17-SEP-2001; 2001FR-00011978.
                                                                                                                                                                                                                                                                                                                                          17-SEP-2002; 2002WO-IB004208
                                                                                                                                                                                                                                                                                                                                                                        27-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tumour suppression related human fukutin oligo SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid sequences associated with tumor suppression, regression apoptosis or virus resistance are useful to diagnose and treat viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-250498/25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence represents an isolated nucleic acid sequence associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          355 CGCAAGGCTGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
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                                                                                                                                                                                                                                                                               MOLECULAR ENGINES LAB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ž
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.0%;
ilarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                Amson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Telerman
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                                                                                                                                                                                                                                                 77
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                                                                                                                                                                                                                                                 Tuijnder M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              degeneration
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The
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CC identifying, quantifying and/or amplifying a nucleic acid, e.g. as one CC component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, collypeptides, vectors containing the nucleic acids, cells containing the cector or antibodies directed against the polypeptides are useful for CC preparation of pharmaceuticals for prevention and/or treatment of viral CC diseases that are characterised by development of tumours or cell CC diseases that are characterised by development of tumours or cell CC diseases. Analysis of the expression of the 17 mer nucleic acids in CC diseases. The polypeptides can also be used to generate antibodies, and CC diseases. The polypeptides can also be used to generate antibodies, and CC both the polypeptide and antibodies are useful as components of protein CC chips. The nucleic acid sequences of the invention can be used in gene charapy. This polynucleotide sequence represents a tumour suppression con the invention can be used in gene contents.
   Sequence 17
BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
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밁
            Ś
                                              Query Match
                                      Local
              355
4
                                      Similarity
CGCAAGGCTGAG
              CGCAAGGCTGAG 366
                               Conservative
                                    100.0%;
                                               2.0%;
                              <u>.</u>.
                                             Score 12;
                                      Pred. No.
                               Mismatches
                                     DB 7; Le
1.2e+05;
                                             Length 17
                             0
                             Gaps
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12-JUN-2003
                             ABT36423
                                                      ABT36423 standard; DNA; 17
(first entry)
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schizophrenia; human fukutin; Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; ge antisense; sense; tumour; cell degeneration; cancer; Alzheimer's protein chip; gene therapy;
ds. tumour suppression; gene chip; disease;

Tumour suppression

related

human fukutin oligo SEQ

占 ö

Homo sapiens

WO2003025175-A2

27-MAR-2003

17-SEP-2002; 2002WO-IB004208.

17-SEP-2001; 2001FR-00011978

Telerman A, Amson Þ Tuijnder M;

(MOLE-) MOLECULAR ENGINES LAB

2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases with tumors and cell degeneration, also related polypeptides, and transfected cells. associated antibodies

Disclosure; Page 273; 720pp; French

RESULT 148
ABT36423/c
ID ABT364
XX ABT364
XX ABT364
XX ABT364
XX Cytost
KW Cytost
KW Schizo
KW Homo s
XX Homo s
XX Homo s
YX H The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for

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RESULT 149
ABT34697
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Best Local
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The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying anucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     schizophrenia;
human fukutin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; ge antisense; sense; tumour; cell degeneration; cancer; Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tumour suppression related human fukutin oligo SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 73; 720pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-SEP-2001; 2001FR-00011978
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 4 A; 5 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein chip; gene therapy; tumour suppression; ds.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ξ,
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1.2e+05;
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vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention
Sequence 17
  BP; 8
  A; 3 C; 4 G; 2 T; 0 U; 0 Other;
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Matches
                                                  Query Match
                                          Local
                 461 ACCATGAAAGAA 472
ហ
                                l Similarity
12; Conserv
ACCATGAAAGAA 16
                                  Conservative
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                                         Score 12;
Pred. No.
                                 Mismatches
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RESULT 150
ADB03447/c
ID ADB03447/c
ID ADB03447/c
ID ADB034
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AC Cytos
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AC Cy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              developmental disorder; ss.
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30-JUL-2002; 2002EP-00016874
                                                                                                                                                                                        EP1281758-A2
                                            02-AUG-2001; 2001US-00922181
(AEOM-) AEOMICA INC
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New zinc finger-containing proteins and nucleic acids, useful manufacturing a medicament for treating or preventing a disord associated with decreased or increased expression or activity MDZ4, MDZ7 or MDZ12, e.g. cancer. disorder ä

Shannon M,

Gu Y,

Nguyen

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WPI; 2003-423107/40

Example 8; SEQ ID NO 4433; 103pp; English.

proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7922.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p21.3 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are The present invention relates to novel human zinc finger-containing

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Best Local S
Matches 12
                                                                                                                                                            The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6g21.3-22.2, MDZ7 is encoded at chromosome 16p1.2 and MDZ12 is encoded at chromosome 16p1.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human MDZ3 scanning oligonucleotide SEQ ID 793.
                                                                                                    useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 8; SEQ ID NO 793; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-423107/40
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                                                                 The present sequence was used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gu Y,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nguyen
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Pred. No.
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RESULT 152
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                                                                                                                                                                                                                                                                                                                                 The present invention relates to novel human zinc finger-containing corporations and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7022.1, MDZ4 is encoded at chromosome 5021.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 sequences are useful in therapy. Cor in manufacturing a medicament for treating or preventing a disorder cassociated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic cacids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic caids can also be used as probes to detect and characterize gross calterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for justices the inventor.
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                                                                                                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 8; SEQ ID NO 794; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-AUG-2001; 2001US-00922181
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                                                   TTTACTGCTGGA 445
                                                                                                                                                                                                                                                                                                             The present sequence was used to
                                                                                                                                                                                                                                                BP; 2 A; 3 C; 4 G; 8 T; 0 U; 0 Other;
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Query Match Best Local

Local Similarity

2.0%;

Score 12; Pred. No.

DB 7; L

Length

Sequence 17

BP; 2 A;

3 C; 5 G; 7 T; 0 U; 0 Other;

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RESULT 154
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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                                                                    20-NOV-2003
                                                                                                                                                                                                  ADA99806 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; SEQ ID NO 312; 103pp; English
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Human MDZ3 scanning oligonucleotide

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Homo sapiens

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RESULT 155
ADB03443/c
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                                                                                    Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7. chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q developmental disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                         Human MDZ7 scanning oligonucleotide SEQ ID 4429.
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                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                          DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Score 12; DB 7; Lenv
%; Pred. No. 1.2e+05;
                                                                                                                chromosome 15q26.1; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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RESULT 156
ADB03445/c
ID ADB034
AC ADB034
AC ADB034
AC ADB034
XX
DT 20-NOV
XX
Cytost
KW Cincon
KW cinc f
KW chromo
KW develo
XX
CS Homo s
XX
OS Homo s
XX
OS FEB
DD 05-FEB
XX
XX
PF 30-JUL
XX
PF 30-JUL
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,
                                                                                                                                                                                                        Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7 chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q
                                                                                                                                                                                                                                                                              Human MDZ7
                                                                                                                                                                                                                                                                                                              20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                 ADB03445
                                                                                                                                                                                                                                                                                                                                                                                 ADB03445 standard; DNA; 17
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                                                   30-JUL-2002; 2002EP-00016874.
                                                                                     05-FEB-2003
                                                                                                                       EP1281758-A2
                                                                                                                                                                                          developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; SEQ ID NO 4429; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-423107/40
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mes 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16
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                                                                                                                                                                                                                                                                            scanning
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 0 A; 8 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                            oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 17;
                                                                                                                                                                                                          15q26.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                          cancer;
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02-AUG-2001; 2001US-00922181

New zinc finger-containing manufacturing a medicament

proteins and nucleic acids, for treating or preventing

B

useful in disorder

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RESULT 157
ADB03446/c
ID ADB034
XX
ADB034
XX
ADB034
XX
ADB034
XX
PT 20-NOV
DT 20-NOV
DT 20-NOV
DT 20-NOV
Cytost
KW Cytost
KW chromo s
KW c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosting or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                         WPI; 2003-423107/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP1281758-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human MDZ7 scanning oligonucleotide SEQ ID 4432.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB03446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADB03446 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 0 A; 8 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New zinc finger-containing proteins and nucleic acids, useful manufacturing a medicament for treating or preventing a disord associated with decreased or increased expression or activity MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                  02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                          05-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 8;
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                                                                                                                                                                                                          (AEOM-) AEOMICA INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 365 AGCCCGAGGGGC 376
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                                                                                                                                                Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                မ်
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gu Y, Nguyen C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                             Y, Nguyen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.2e+05;
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disorder
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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RESULT 158
ADA99807
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     associated v
                     New zinc finger-containing proteins and nucleic acids, useful manufacturing a medicament for treating or preventing a disornassociated with decreased or increased expression or activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
                                                                                                                         WPI; 2003-423107/40.
                                                                                                                                                                     Shannon M,
                                                                                                                                                                                                                                                                  02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                30-JUL-2002; 2002EP-00016874
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human MDZ3 scanning oligonucleotide SEQ ID 796.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to novel human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  365 AGCCCGAGGGGC 376
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                                                                                                                                                                                                                   AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       with decreased or MDZ12, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 4432; 103pp; English.
or MDZ12, e.g. cancer.
                                                                                                                                                                   Gu Y, Nguyen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 1 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                        d nucleic acids, useful in g or preventing a disorder expression or activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 7; Lo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        zinc finger-containing
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3: encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-

or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 5g21.3-22.2, MDZ7 is encoded at chromosome fighl: 2 and MDZ12 is encoded at chromosome 15g1.2 and MDZ12 is encoded at chromosome 15g26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy,

Example 8; SEQ ID NO 798; 103pp; English

Example 8;

SEQ ID NO 796; 103pp; English.

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RESULT 159
ADA99809
ID ADA998
XX ADA998
XX ADA998
XX ADA998
XX Cycon
DE Human
XX Cycon
Chromo
KW Chromo
KW Chromo
CX Homo s
XX EP1281
XX BP1281
XX BP1382
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                 The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy. Or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The probes are useful as therapeutic agents for gene therapy or as
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                                                                                                                                                                                                                                                                                                                                                        New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                             The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are
                                                                useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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ARESULT 162
ADB03444/c
ID ADB03444/c
ID ADB03444/c
AC ADB034
AXX
AXX ADB034
AXX
DTT 20-NON
XXX CYtos
KW Cytos
KW Cytos
KW Chron
CKW Chro
RESULT 163
ADA99808
ID ADA998
XX
AC ADA998
XC
AC ADA998
XX
DT 20-NOV
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7922.1, MDZ4 is encoded at chromosome 6p21.3-22.2, CMDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome (15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy. Co or in manufacturing a medicament for treating or preventing a disorder cassociated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic caids and proteins are also useful for diagnosing or monitoring a disease CC caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic caids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The CC useful in constructing microarrays for measuring gene expression. The CC vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New zinc finger-containing proteins and nucleic acids, useful manufacturing a medicament for treating or preventing a disord associated with decreased or increased expression or activity MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB03444 standard; DNA;
                                   20-NOV-2003
                                                                                                                                                                    ADA99808 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JUL-2002; 2002EP-00016874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP1281758-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human MDZ7 scanning oligonucleotide SEQ ID 4430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                         15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8; SEQ ID NO 4430; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                        AGCCCGAGGGGC 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 0 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                   (first entry)
                                                                                                                                                                    DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12; DB 7; L
Pred. No. 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17;
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ABEGIJT 164
ABEG1994/c
ID ABEG19
XX ABEG19
XX ABEG19
XX ABEG19
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XX Human
DE Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 Is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p12.2 and MDZ12 is encoded at chromosome 25p26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, 25p27 contacted with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic 25p27 contacted with decreased of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic 25p27 contacted by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic 25p2 cancer on the 25p2 contacted by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are 25p2 useful in constructing microsarrays for measuring gene expression. The 25p2 proteins are useful as therapeutic agents for gene therapy or as 25p2 contacted and character the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p11,3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 2 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-AUG-2001; 2001US-00922181.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    developmental disorder; ss.
                                                            Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                 ABZ61994 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human MDZ3 scanning oligonucleotide SEQ
                                                                                                                                           Human H-Ras
                                                                                                                                                                                  21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 8; SEQ ID NO 797; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JUL-2002; 2002EP-00016874.
                                                                                                                                                                                                                                                                                                                                                                                                           434
                                                                                                                                                                                                                                                                                                                                                                   N
                                                                                                                                                                                                                                                                                                                                                                   TTTACTGCTGGA 13
                                                                                                                                                                                                                                                                                                                                                                                                        TTTACTGCTGGA 445
                                                                                                                                           DNAzyme target #785
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.2e+05
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RESULT 165
ABZ61385
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Best Local S
Matches 12
                                   29-MAY-2001;
06-JUN-2001;
10-SEP-2001;
                                                                                                                                                                                                                                                             Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human references.
                                                                                                          29-MAY-2002; 2002WO-US016840
                                                                                                                                            05-DEC-2002.
                                                                                                                                                                             WO200297114-A2
                                                                                                                                                                                                                                             anti-rheumatic; cancer; AIDS; ss
                                                                                                                                                                                                                                                                                                                 Human H-Ras DNAzyme target #176.
                                                                                                                                                                                                                                                                                                                                                    21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ61385 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ribozymes of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-MAY-2002; 2002WO-US016840
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     (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel short interfering RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CCCCAACCTCAG 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 126; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 2 A; 3 C; 8 G; 0 T; 4 U; 0
                                   2001US-0294140P.
2001US-0296249P.
2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                      RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 166
ABZ61383
ID ABZ61383
XX ABZ613
XX ABZ613
XX Human
XX Human
XX Human
XX Human
XX Homo E
XX MO2002
XX NOVELL
PT HER2,
XX NOVELL
PT HER2,
XX Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                 Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                          29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                   Claim 58; Page 114; 185pp;
                                                                                                                        WPI; 2003-140484/13.
                                                                                                                                                                                                                                                                                               29-MAY-2002; 2002WO-US016840
                                                                                                                                                                                                                                                                                                                                 05-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human H-Ras DNAzyme target #174.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ61383 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 0 A; 4 C; 12
                                                                                                                                                                                                                                                                                                                                                                  WO200297114-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 58; Page 114; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-140484/13
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                                                                                                                                                                                          (RIBO-) RIBOZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                373 есестесессе 384
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                            PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.0%;
91.7%;
                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12;
Pred. No.
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65551, ABZ66520 - ABZ66524, ABZ66530 - ABZ66530 - ABZ66530.
Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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RESULT 167
ABZ61791/c
ID ABZ617
XX ABZ617
XX ABZ617
XX ABZ617
XX Human
DE Human
XX Human
XX Homo 8
PN W02002
XX Homo 8
PN W02002
XX 29-MAY
XX 29-MAY
PR 06-JUN
PR 10-SEI
XX HOVel
PT HER2,
XX WPI;
YX VICE
PT HER2,
XX Claim
XX Claim
XX Claim
CC Also
CC Shown
CC ABZ66
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression of a nucleic acid molecule, that modulates human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancratic cancer, and HIV infection, and AIDS. The sequences shown in ABZ5989 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ6530 - ABZ6585 represent substrate/target sequences for the human ribozymes of the invention
                 The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAY-2001; 2001US-0294140P
06-JUN-2001; 2001US-0296249P
10-SEP-2001; 2001US-0318471P
                                                                                                                                                                                                                                                                                                   treating cancer, mod
HER2, K-Ras, H-Ras,
                                                                                                                                                                                                                                                                                                                          Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
                                                                                                                                                                                                                                                           Claim 58; Page 122; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-140484/13
                                                                                                                                                                                                                                                                                                                                                                                                                                     Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAY-2002; 2002WO-US016840
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human H-Ras DNAzyme target #582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ61791 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          372 ееестесес звз
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  ABZ66585 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 0 A; 5 C; 10 G; 0 T; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lates to a novel short interfering RNA (siRNA) nucleic an enzymatic nucleic acid molecule, that modulates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.0%;
91.7%;
                                                                                                                                                                                                                                                                                                        N-Ras,
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Pred. No.
    substrate/target
                                                                                                                                                                                                                                                                                                        human
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                                                                                                                                                                                                                                                                                                        deficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2e+05;
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    sequences
                                                                                                                                                                                                                                                                                                        sequences.
      for the
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    human
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CTGTGCCCGACA 278

Query Match Best Local : Matches

l Similarity
12; Conserv

Conservative

0;

Mismatches

Indels

0

Gaps

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100.0%;

2.0%;

Score 12; DB 7; I Pred. No. 1.2e+05;

Length 17;

Sequence 17

BP; 3 A; 4 C; 6 G; 0 T; 4 U; 0 Other;

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RESULT 168
ABZ61790/c
ID ABZ617
XX ABZ617
XX ABZ617
XX ABZ617
XX Human
XX Human
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Best Local S
                                                                                     acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human in ABZ6585 represent substrate/target sequences for the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ61790 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAY-2001; 2001US-0294140P
06-JUN-2001; 2001US-0296249P
10-SEP-2001; 2001US-0318471P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            anti-rheumatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human H-Ras
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-MAR-2003
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                                                                     ribozymes of
                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 58; Page 122; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-140484/13
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12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer; AIDS; ss.
                                                                     invention
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Pred. No.
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. 1.2e+05;
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RESULT 170
ACC63317
ID ACC653
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ACC64654
                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                             ACC68866), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonuclectides are useful as (1) as probes and primer received, e.g. as one component of a gene chip; in vitro as (anti)sense reagents; and (2) for production of recombinant polypeptides. The oligonuclectides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-SEP-2001; 2001FR-00011979
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUL-2003
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Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; mur tumour suppression; tumour reversion; apoptosis; virus resistance;
                                          Murine oligonucleotide associated with tumour supression,
                                                                        01-JUL-2003
                                                                                                                                                                                                                                                                                                                     Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 253; 738pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAR-2003
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                                                                                                                                ACC65317 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to murine oligonucleotides (ACC62754-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2003-333167/31.
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                                                                                                                                                                                                    TGAGACCTACCT 16
                                                                                                                                                                                                                               TGAGACCTACCT 86
                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                     BP; 5 A; 5 C; 3 G; 4 T; 0 U; 0
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                                                                       (first entry)
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                                                                                                                                DNA; 17
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                                                                                                                                                                                                                                                                                                                     Other;
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                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                       ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, of a quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip; in vitro as (ant.) sense reagents, and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases the characterised by development of tumours or cell degeneration, are difficulty of the properties of tumours or cell degeneration,
 ACC64816
                                                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               viral disease;
                                                                                                                                                                                                                            specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to murine oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 330; 738pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Telerman A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    schizophrenia;
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                                                                                                                                    l Similarity
12; Conserv
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                                                                                                       CTGGACTTTGGT 141
standard; DNA; 17
                                                                           CIGGACTITGGT 17
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                                                                                                                                   larity 100.0%; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amson R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tumour; cell degeneration; cancer; Alzheimer's disease,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour suppression; viral disease; tumou schizophrenia; ss.
                                                                                                                     17-SEP-2001; 2001FR-00011979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Murine oligonucleotide associated with tumour supression,
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Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip; in virro as (anti)sense reagents, and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
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04-DEC-2003
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                                                                                                                                    usetul e.g.
polypeptide
                                                                                                                                                          New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also relat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss; primer; probe; tumour suppression; tumour reversion; apoptosis;
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                                                                           Disclosure; Page 155; 771pp;
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                                                                                                                                                                                                                                                                                                                                                       (MOLE-) MOLECULAR ENGINES LAB
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(first en
                                                                                                                                      and antibodies.
                                                                                                                                                                                                                                                                                                 Amson R,
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                                                                                                                                                                                                                                                                                              Tuijnder
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Pred. No.
                                                                                 French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              animals; Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        associated nucleotide
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o. 1.2e+05;
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                                                                                                                                                                also related
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The invention fragments of a

n relates at least

15 15

the isolation of 6327 nucleotide consecutive nucleotides of these

sequences, nucleotides,

The invention relates to the isolation of 6327 nucleotide sequences, fragments of at least 15 consecutive nucleotides of these nucleotides, sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the

Disclosure;

Page 443; 771pp;

French

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cc sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with ct the nucleotides. Fine nucleotides are used as probes or primers for detecting, cidentifying, quantifying and/or amplifying nucleic acids, as in vitro complete and antisense sequences, of nucleotides involved in tumour customers and antisense sequences, of nucleotides involved in tumour customers and to prepare transgenic animals, as an ecombinant polypeptides, and to prepare transgenic animals, as cells containing the vectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment cor real degeneration (e.g. Alzheimer's disease or schizophrenia).

Cc Analysis of the expression of the nucleotides can be used for diagnosis and ob used to screen for their specific interactive molecules, con potentially useful for treating diseases associated with abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                            New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                                                                                                                                                                                                                       17-SEP-2001; 2001FR-00011981
                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                              primer; probe; tumour suppression; tumour reversion; apoptosis;
virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;
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04-DEC-2003
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                                                                                                                                                                     WPI; 2003-441574/41.
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CC analysis of the expression of the nucleotides can be used for diagnosis CC and/or prognosis of these diseases. The nucleotides and polypeptides can cC calso be used to screen for their specific interactive molecules.
CC potentially useful for treating diseases associated with abnormal cover of the nucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                    New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
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                                                                                                                                                                                                                                                           Telerman A,
                                                                                                                                                                                                                                                                                                                     17-SEP-2001; 2001FR-00011981.
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                           Amson R,
                                                                                                                                                                                                                                                                                                                                                 2002WO-IB004219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                          ENGINES LAB
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               schizophrenia;
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The invention relates to the isolation of 6327 nucleotide sequences, fragments of at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro sense and antisense sequences, of nucleotides involved in tumour suppression or reversion, apoptosis and or viral resistance, to produce

Query Match Best Local Similarity

2.0%; 100.0%; 2 C;

Score 12; Pred. No.

DB 2; L

Length 18;

Sequence

18

BP; 4

2

w <u>.</u>

9 T;

0 U;

0

Other;

Disclosure;

Page 687; 771pp; French

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RESULT 175
AAQO6031/c
ID AAQO60
XX AAQO60
XX AAQO60
XX Detect
XX Detect
XX Detect
XX SS.
XX USN746
XX Detect
PF 24-JAN
XX USN746
XX USN746
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PF 26-JAN
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Best Local S
Matches 12
                                                granted patent numbers. For further information please web site at www.derwent.com/dwpi/updates/ntis_us.html.)
                                                                           The labelled probes whose SQs are given in AAQ06030-2 are used for detecting amplified VP1 amino terminal genome region HAV cDNA. The met of the invention provides a rapid, sensitive and specific test for detecting the presence of HAV in infected or contaminated samples of e.g., stools, serum, water or foodstuffs such as shellfish. (Note: Revised entry submitted to correct the patent number format of US covernment cowned NTIS applications to prevent clashes with ongoing US granted patent numbers. For further information please visit the Derweigh the patent numbers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         recombinant polypeptides, and to prepare transgenic animals, as experimental models. The nucleotides (also vectors containing them and cells containing the vectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment of viral infections or diseases characterized by development of tumours or cell degeneration (e.g. Alzheimer's disease or schizophrenia) analysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can also be useful for treating diseases associated with abnormal expression of the nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Detection of hepatitis A virus in synthesising cDNA, amplifying the
                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 6; 28pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1990-304798/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence of VP1 (-) sense probe terminal genome region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-DEC-2001
08-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JAN-1990;
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US SEC OF COMMERCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90US-00469143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A virus;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   viral protein 1; polymerase chain
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                                                                                                                                                                                                                                                                                                                                                                                                                                         samples cDNA by
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    by extracting
PCR and detection

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RNA,
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RESULT 176
AAQ26186/c
                                                                 RESULT 177
AAQ26149
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                                                                                                                                                                                                                                                                                                                 The sequence is that of the hybridising region of tailed probe DRB82 for use in a method for determining HLA-DR beta sub-type in a nucleic acid sample. The method allows specific nucleic acid sequences of the second exon of HLA-DR beta genes to be amplified then probed for identification of polymorphic sequences. The amplified DNA is useful for typing homozgous or heterozygous samples from a variety of sources and for detecting allelic variants not distinguishable by serological methods. The typing system can be used in a reverse dot blot format which is simple and rapid to perform, produces detectable signals in minutes and can be utilised in tissue typing, determination of individual identity and identifying disease susceptible individuals. See also AAQ26092-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Method for determining HLA-DR beta sub-type in DNA sample - comprises amplification and hybridisation with probes and primers, useful in tissue
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04-JAN-1993
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                   AAQ26149;
                                                  AAQ26149 standard; DNA; 18
                                                                                                                                                                                                                                                                     Sequence 18 BP; 2 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Erlich HA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-DEC-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tissue typing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HLA-DR beta sub-type tailed probe DRB82 hybridising region.
                                                                                                                                                                                                                                                                                                      Q26367. (Updated on 25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example; Page 39; 90pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HOFF ) HOFFMANN LA ROCHE & CO AG F.
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                                                                                                                                                          244 ACCTCCTGGAGC 255
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(first en
                                                                                                                                                                                                     Conservative
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                                                  βP
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Pred. No.
                                                                                                                                                                                                       Mismatches
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io. 1.2e+05;
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RESULT 178
AAQ38340
ID AAQ383
XX AAQ383
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DT 25-MAR
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Erlich HA,
Apple RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence is that of the hybridising region of tailed probe DRB44 for use in a method for determining HLA-DR beta sub-type in a nucleic acid sample. The method allows specific nucleic acid sequences of the second exon of HLA-DR beta genes to be amplified then probed for identification of polymorphic sequences. The amplified DNA is useful for typing homozogous or heterozygous samples from a variety of sources and for detecting allelic variants not distinguishable by serological methods. The typing system can be used in a reverse dot blot format which is simple and rapid to perform, produces detectable signals in minutes and can be utilised in tissue typing, determination of individual identity and identifying disease susceptible individuals. See also AAQ26092-Q26367. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                   EcoRI primer
                                                                                                                                                                                    25-MAR-2003
15-JUL-1993
                                                                                                                                                                                                                                                                     AAQ38340 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 3 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Method for determining HLA-DR beta sub-type in DNA sample - amplification and hybridisation with probes and primers, use
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04-JAN-1993
                                  EP534858-A1
                                                                   Synthetic.
                                                                                                                  Restriction
                                                                                                                                                                                                                                    AAQ38340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 38; 90pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-DEC-1991;
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                                                                                                  selective restriction
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                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
12; Conserv
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                                                                                                              fragment;
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(first entry)
                                                                                                                                                                                  (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                       2.0%;
                                                                                                 PCR; RFLP; forensic typing; fragment amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bugawan T,
                                                                                                                                                   corn
                                                                                                                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                         <u>,</u>
                                                                                                                                                   DNA with multiple restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Griffith
                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2; Le
. 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 18
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                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                  Zea mays; ss; labelled;
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31-MAR-1993

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RESULT 179
AAQ34757
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The DNA from two corn inbred lines was restricted with TaqI and one of AseI, PstI, EcoRI or Sse8387-I and tagged with the appropriate adapters. This DNA was used as template in PCR reactions using one of the labelled TaqI primers, 1-4 and one of PstI-primers 1-4, AseI primers 1-4, CoRI-primers 1-4 or Sse8387-I primers 1-4, each primer contg. three selective nucleotides at the 3' end. A total of 128 PCRs was performed and the reaction prods. analysed. All primer combinations gave DNA fingerprints of 50-100 bands per lane, except for the combination SseI/TaqI, which gave only 10-15 bands per lane. The method can be used to identify preselected DNA fragments which can be polymorphic, i.e. RFLBs. It is a superior method for multiplex PCR. The method may also be used to detect evaluating genetic distances and characterising the plant. See also called the second plant or animal varieties, species etc, or for evaluating genetic distances and characterising the plant. See also called the second plant and AAQ38907-16. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                        AAQ34757 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Selective controlled restriction fragment amplification of DNA - used for gene analysis e.g. DNA fingerprinting, restriction fragment length
                                                                                                                                 06-JUL-1992;
                                                                                                                                                              21-JAN-1993
                                                                                                                                                                                            WO9301313-A1
                                                                                                                                                                                                                                                                                                                 25-MAR-2003
03-JUN-1993
                                                                                                                                                                                                                                                                                                                                                            AAQ34757
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polymorphisms, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1993-102942/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zabeau M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-SEP-1992;
              WPI; 1993-045514/05
                                          Cardy DLN,
                                                                                                   05-JUL-1991;
                                                                                                                                                                                                                                                     Human; p53; anti-oncogene; assay; ss
                                                                                                                                                                                                                                                                                   Activator oligonucleotide probe p53(A)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (KEYG-) KEYGENE NV.
                                                                      (CYTO-) CYTOCELL LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 TTACCAATTCAA 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                          Delnatte SYJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 5 A; 6 C; 2 G; 5 T; 0 U; 0 Other;
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(first entry)
                                                                                                   91GB-00014525
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                                                                                                                                  92WO-GB001229
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100.0%; Pred. No. 1.2e+0
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RESULT 180
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Best Local S
Matches 12
                                                            Query Match
Best Local &
                                                 Matches
                                                                                                                        The blocker oligonucleotide p53(B) is complementary to its corresp. activator probe p53(A). The oligonucleotide was synthesised with a 5' bromodeoxyuridine nucleotide and may be used in an assay for the detection of the human p53 anti-oncogene in human DNA. See also AAQ34 62. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                           Homogeneous assay for nucleic acid sequences - obtd. by modulating enzyme activity, by hybridisation of derived nucleic acid probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ34759;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 4 A; 6 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                  Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; p53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blocker oligonucleotide p53(B)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003
03-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The activator oligonucleotide probe p53(A) may be used in an the detection of the human p53 anti-oncogene in human DNA. Se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 27; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homogeneous assay for nucleic acid sequences - obtd. by modulating enzyme activity, by hybridisation of derived nucleic acid probes.
                                                                                                                                                                                                  Example 2; Page 27; 51pp; English.
                                                                                                                                                                                                                                                                 WPI; 1993-045514/05.
                                                                                                                                                                                                                                                                                                                                            05-JUL-1991;
                                                                                                                                                                                                                                                                                                                                                                     06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                      W09301313-A1
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                                                                                                                                                                                                                                                                                          Cardy DLN, Delnatte SYJ;
                                                                                                                                                                                                                                                                                                                   (CYTO-) CYTOCELL LTD
                                                                                                                                                                                                                                                                                                                                                                                              21-JAN-1993.
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                        9,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     il Similarity
12; Conserv
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                                                 12;
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                                                             Similarity
                        ATGAACCGGAGG 27
 ATGAACCGGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATGAACCGGAGG 15
                                                 Conservative
                                                                                                BP; 1 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anti-oncogene; assay; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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(first entry)
                                                                                                                                                                                                                                                                                                                                            91GB-00014525
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.0%;
                                                            2.0%; Score 12; DB 2; L
100.0%; Pred. No. 1.2e+05;
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Pred. No.
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                                                 Mismatches
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                                                                         Length 18;
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                                                 Indels
                                                                                                                                        DNA. See also AAQ34750-
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                                               Gaps
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RESULT 181 AAV39342/c

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RESULT 182
AAZ21414
ID AAZ214
XX
AC AAZ214
XX
DT 02-DEC
XX
DE Human
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                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a PCR primer for use in a method of the cinvention for determining the genetic predisposition to cancer in an individual by detecting hands mutations in a sample. hRAD54 is a gene thought to be present in tumours that display allelic imbalance at Ip32, the chromosomal band identified as one of four minimal regions of chromosome 1 deletion in breast carcinomas hRAD54 is useful for production of proteins, inter alia, that have been identified as novel hRAD54 by homology between the amino acid sequence given in AAM62186 and known amino acid sequences such as yeast RAD54. hRAD54 proteins are used in the treatment of cancer, including Xeroderma Pigmentosum and Bloom syndromes and Sequences and X-linked mental retardation with alphacthalisasemia syndrome and breast cancer. hRAD54 polymucleotides are also useful for detecting complementary nucleotides for use as a diagnostic agent, especially useful for diagnosis of disease or susceptibility to disease hands a protein a susceptibility to disease hands a protein a protein a protein and such as the protein are also useful for detecting complementary nucleotides for use as a diagnostic disease hands are also agent, especially useful for diagnosis of disease or susceptibility to disease the protein are also as the protein and the protein are also as the protein and the propen and the protein and the protein and the protein and the prote
                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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                                         02-DEC-1999
                                                                                                                      AAZ21414 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 3 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 18; Page 50; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Werner's syndrome; ATR-X; X-linked mental retardation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human RAD54 mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV39342 standard;
Human MEK2 antisense oligonucleotide SEQ ID NO:17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SMIK ) SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                               proteins
                                                                                                                                                                                                                                                                487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hRAD54 DNA and polypeptide -
                                                                                                                                                                                                                           12
                                                                                                                                                                                                                                                                                                        12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RAD54; hRAD54; cancer; xeroderma pigmentosum; Bloom syndrome;
's syndrome; ATR-X; diagnosis; detection; SNF2 superfamily;
                                                                                                                                                                                                                                                              GAAGGGCTGCAT 498
                                                                                                                                                                                                                         GAAGGGCTGCAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                           hRAD54 polynucleotide, proteins, agonists and antagonists which ns are useful in gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fishel RA,
                                                                                                                                                                                                                                                                                                      Conservative
                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96US-0030676P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   retardation with alpha-thalassemia syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA;
                                                                                                                                                                                                                                                                                                                         100.0%;
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                                                                                                                                                                                                                                                                                                                                             2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rasio D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                           Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer SEQ ID NO:50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Robbins DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and agonists, antibodies, antagonists,
                                                                                                                                                                                                                                                                                                                                             DB 2;
                                                                                                                                                                                                                                                                                                                           1.2e+05;
                                                                                                                                                                                                                                                                                                                                           Length 18
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UX AX FX BX BX S S X R
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes antisense oligonucleotides (asMEK2) of the human MEK2 dual specificity kinase gene. The MEK2 gene (also called MAPKK2, mpk2, MAPK/ERK kinase 2, prmmk2 and erk activator kinase) represents a convergent target for the regulation of a range of cellular processes including proliferation, differentiation and development asMEK2 may be used to inhibit the expression of MEK2 genes in vivo and/or in vitro. MEK2 has been shown to be over expressed in some tumour cells. Therefore, asMEK2 may be administered to a patient to treat hepatocellular carcinomas and breast carcinomas. asMEK2 may also be used as a diagnostic tool to specifically inhibit the expression of MEK2 genes. The role of the inhibited genes in cellular pathways may then be evaluated. They may also be used as probes to detect sequences encoding MEK2 or as primers for the amplification of those sequences. AAZ21406 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mitogen activated protein kinase; expression; modulation; antisense; diagnosis; hepatocellular carcinoma; breast cancer; proliferation; differentiation; development; detection; probe; primer; tumour;
                                                                                                G-alpha-13;
                                                                                                                                                                                                                                                              AAZ31807 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 1 A; 3 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense oligonucleotides of the MEK2 kinase gene, useful in diagnostic protocols in vitro and for inhibiting the expression of MEK2 in vivo for the treatment of human hepatocellular carcinomas and breast carcinomas.
US5981732-A
                                                          Synthetic
                                                                                                                                       Human G-alpha-13 antisense inhibitor ISIS# 20756.
                                                                                                                                                                                 24-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ21443 represent specifically claimed asMEK2's from the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; MEK2; MAP kinase; MAPK/ERK kinase; erk activator kinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                        130 CTGGACTTTGGT 141
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                                                                                                                                                                                                                                                                                                                                                                CTGGACTTTGGT 17
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                                                                                                  human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                    inhibitor;
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Pred. No.
                                                                                                cancer; antisense compound; therapy; ss
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Best Local S
Matches 12
                                                                      New polynucleotide from Escherichia coli encoding 6-phosphogluconate dehydrogenase and polymeric sequences in the nucleotide for detecting strains of E. coli for diagnosis and food screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents an antisense inhibitor of the invention, and inhibits the expression of the human G-alpha-13 protein. The antisense compounds of the invention are of 8 to 30 nucleobases in length, that inhibits the expression of the human G-alpha-13. The antisense compound is useful for treating an animal, particularly humans, having or being
                                                                                                                                                                                     WPI; 2000-431304/37.
                                                                                                                                                                                                                                                                                                                                               08-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                   08-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Escherichia coli 055:H7; gnd; 6-phosphogluconate dehydrogenase; 6-PGD; gastrointestinal infection; pathogen; 0157 lipopolysaccharide; haemolytic uraemic syndrome; HUS; haemolytic anaemia; thrombocytopenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA52916;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200034484-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Escherichia coli 055:H7 gnd gene PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA52916 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 5 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense compound inhibiting expression of human G-alpha-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              to a disease or condition associated with the expression of G-alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7
                                                                                                                                                                                                                                                                                         CHILDREN'S HOSPITAL & REGIONAL MEDICAL.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 18
                                                                                                    detecting
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RESULT 185
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is a PCR primer used to amplify DNA beyond the 5'CC and 3' termini of the E. coli 055:H7 gnd gene from total genomic DNA of E. coli 055:H7 strain. The resulting sequence data was used to design a CC primer pair to amplify the E. coli 055 gnd allele. The gnd gene, which can be used to amplify the E. coli 055 gnd allele. The gnd gene, which coli content strains of E. coli. Several polymorphisms have been found that CC an be used to identify the presence of a particular strain of E. coli cand/or to differentiate one strain from another. A substitution of an including mild distrains of 0157:H7 and coli can be used to differentiate thighly pathogenic strains of 0157:H7 and coli of the coli of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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            This is a human G-alpha-12 antisense nucleotide sequence. G-alpha-12 is member of the G12/13 subfamily of G-proteins. The primary function of G-alpha-12 is in cell differentiation and growth. The invention relates to antisense compounds which are 8-30 nucleotides long (see AAE57666-Z57746). The antisense molecules are targeted to the human G-alpha-12 nucleic acid molecule, and inhibit the expression of G-alpha-12. The molecules preferably have a modified internucleotide linkage, and at least one modified sugar molety. The compounds target different regions of the human G-alpha-12 RNA. The expression of human G-alpha 12 is inhibited by contacting human cells or tissues in vitro with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G-alpha-12 inhibitor; antisense
                                                                                                                                                                                                                                                                                                Example 15; Col 38; 36pp; English.
                                                                                                                                                                                                                                                                                                                                               Antisense inhibition of human G-alpha-12 expression
                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-095920/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human G-alpha-12 antisense inhibitor ISIS# 20661.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 Similarity
12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CCGCTACATOTC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease and to test for food or water contamination
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%; Pred. No. 1.2
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell differentiation; cancer;
20661.
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RESULT 186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel oligonucleotide (I) comprising an adapter sequence and part of the target sequence of a restriction endoublease, and which has 1-10 selected nucleotides immediately adjacent to the 3' end of the target sequence. The products of the invention are used to tag restriction fragments which are to be amplified by the polymerase chain reaction (PCR). This technique may be used in the detection of restriction fragment polymorphisms (RFPs), including length polymorphisms. The products can also be used for genetic analysis, such as for the forenaic typing of humans and the detection of the inheritance of determined traits in animals or plants and to monitor several diseases at once. The oligonucleotides and kits may also be used to identify species, races or varieties of animals or plants. The new adapters, oligonucleotides and methods for using them are more sensitive for the terraction fragment polymorphisms because not only differences in the terraction fragment polymorphisms because not only differences in the terraction fragment polymorphisms because not only differences in the terractions fragment polymorphisms because not only differences.
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        function of nucleic acid molecules encoding G-alpha-12, ultimately modulating the amount of G-alpha-12 produced. The antisense compounds can be utilized for diagnostics, therapeutics, prophylaxis and as research agents and kits. They may be useful in the treatment of cancer, and metastatic growth
in the target sites of the restriction endonuclease are detected as with prior art methods and oligonucleotides but also differences in the adjacent nucleotide sequences within the selective PCR primers. Multiplex PCR may only be used to monitor 5-8 different traits simultaneously and compromise conditions have to be established to allow all primer pairs to yield detectable products. In addition there are strong differences in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-SEP-1991;
24-SEP-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                           genetic diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                          New oligonuclectides, useful for tagging restriction fragments
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  corn; primer;

    mais restriction fragment amplification EcoRI-primer-3.

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07-APR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ44508;
                                                                                                                                                                                                                                                                                                                                                                                       Example 4; Page 18; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ44508 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KEYG-) KEYGENE NV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vos P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 forensic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 4 A; 6 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92EP-00402629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 91EP-00402542
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              restriction fragment polymorphism; multiplex PCR;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 T; 0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 18;
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RESULT 188
AAC66835
ID AAC668
XX
AC AAC668

AAC66835 standard; DNA; 18

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AAC66835;

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RESULT 187
ABA92997
ID ABA929
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                                                                                                                        Query Match
Best Local Similarity
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Best Local
                                                                                                                                                                                                                                                                                The present invention describes angiogenin inhibitors. Angiogenin inhibitors can be used in the treatment of angiogenesis. The presequence represents an oligonucleotide which is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the efficiency of amplification of different fragments and products of certain primer pairs are not detectable with multiplex PCR. In contrass using the new techniques, all the primers have a substantial part of their nucleotide sequence in common and by selecting Amplified Fragmen Length Polymorphisms, the DNA markers are amplified with equal efficiency. AAZ44475-Z44526 represent primers used to illustrate the method of the invention. (Updated on 06-AUG-2003 to correct OS field.)
                                                                                                                                                                                                        Sequence 18 BP; 2 A; 2 C; 5 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                           exemplification of
                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 6; 18pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Angiogenin inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-254109/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chae CB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GREC )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KR99021167-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Angiogenin inhibitor; angiogenesis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Angiogenin inhibitor related oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA92907;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA92907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 5 A; 6 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-AUG-1997;
                                                547 CTGTATGAGGTT 558
                                                                                                     12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      36 TTACCAATTCAA 47
  N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 UNIV POHANG SCI & TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
CIGIATGAGGIT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lee
                                                                                                                                                                                                                                                      the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                             100.0%;
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                                                                                                                                                        2.0%;
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                                                                                                     0; Mismatches
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                                                                                                                                Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 12;
Pred. No.
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1.2e+05;
                                                                                                                                                        B
                                                                                                                             1.2e+05;
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                                                                                                                                                   Length 18
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                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                        present
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                                                                                                Gaps
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RESULT 189
AAH47521
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                       be involved in signal transduction in the cell, and to have binding activity for other telomere-associated proteins. It is possible that it plays a role in the regulation of telomere length, thus affecting the replicative ability of the cell. The protein is useful for ribosylating target proteins, for determining tankyrase II binding activity in a sample, and for modulating telomere length in a cell. The present sequence is a PCR primer used in the isolation of the tankyrase II codir
                                                                                                                                                                30-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to the isolation of the protein and codin sequences of human tankyrase II. The tankyrase II protein is thought to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel mammalian Tankyrase II polypeptide and the polynucleotide encoding the polypeptide useful for modulating or maintaining telomere length, replicative capacity, apoptosis, chromosome packing or gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
                         US1973-H.
                                                                                   anti-arthritic; cytostatic; anti-Parkinsonian; neuroprotective;
nootropic; cancer; apoptosis; Parkinson's disease; Alzheimer's disease;
                                                                                                                                       Human MMP-8 cDNA amplifying sense primer E.
                                                                                                                                                                                         AAH47521;
                                                                                                                                                                                                                   AAH47521 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 9; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Morin GB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-OCT-2000
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                                               Homo sapiens
                                                                        Huntington's
                                                                                                               MMP-8alt; MMP-8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-APR-1999;
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                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                                                         246 CTCCTGGAGCCC 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tankyrase
                                                                                                                                                                                                                                                                                                                                   12;
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                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                    BP; 3 A; 7 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                (first entry)
                                                                          disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
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99US-0129123P.
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                                                                                                             matrix metalloproteinase; neutrophil collagenase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      II; telomere
                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                               2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer gt11-5'
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                                                                                                                                                                                                                   18
                                                                                                                                                                                                                   BP
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                                                                                                                                                                                                                                                                                                                                             Score 12; DB 3; L
Pred. No. 1.2e+05;
                                                                       PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   length; signal transduction; PCR primer;
                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                          Length 18;
                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                 0
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RESULT 190
AAC99241
ID AAC992
XX AAC992
XX AAC992
XX Probe
XX Probe;
XX Probe;
XX Probe;
XX Probe;
XX JP2000
XX JP20
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA. The MMP-Balt polypeptide can be expressed by standard recombinant methodology. The polynucleotides and polypeptides may be used as research reagents and materials for the discovery of treatments and diagnostics to human disease, and as targets for identifying modulators. Inhibitors of MMP-Balt polynucleotide or polypeptide expression may be used to treat and/or prevent arthritis, cancer and cancer metastasis, and diseases caused by cellular apoptosis including Parkinson's disease, Alzheimer's disease and Huntington's disease. Sequences AM447517-21 represent PCR
                                                                                                     A preparation of a probe-combined substrate, a probe array, dete a target substance, specification of the base sequence of a sing stranded nucleic acid in a sample, and determination of a target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probe sequence used in probe array SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC99241 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 7 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New MMP-Balt polynucleotides and polypeptides useful as research reagents and materials for discovering treatments and diagnostics to human disease, or as targets for identifying inhibitors of MMP-Balt expression.
                                                                                   substance in a sample.
                                                                                                                                                                                                                           WPI; 2001-027424/04.
                                                                                                                                                                                                                                                                                                                                               28-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JP2000270896-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Probe; probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAC99241;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primers for cDNA and genomic DNA cloning of MMP-8alt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to human MMP-8alt polypeptide and polynucleotides. MMP-8alt is a splice variant of the MMP-8 (matrix metalloproteinase)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hu S;
                                                                                                                                                                                                                                                                                                                                                                                                        28-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-OCT-1998;
                                                                                                                                                                                                                                                                                       (CANO ) CANON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94 GTAGTGAAGAGG 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       array; probe-combined substrate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                               99JP-00019915
                                                                                                                                                                                                                                                                                                                                                                                                        99JP-00019915.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.2e+05;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       83
                                                                                                                                        y, detection
a single-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>,,</u>
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Example 3;

Page

11;

20pp; Japanese

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RESULT 192
AAC91122
ID AAC911
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AAC99283/c
ID AAC997
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                                                                                                       Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                 A preparation of a probe-combined substrate, a probe array, dete
a target substance, specification of the base sequence of a sing
stranded nucleic acid in a sample, and determination of a target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to a probe-combined substrate, a probe array, and a method for the detection of a target substance in a sample. The probe array can be used for detecting a target substance with high reliability. Sequences AAC99241 - AAC99305 represent probes used in an array in an
                                                                                                                                                                                     This invention relates to a probe-combined substrate, a probe array, and a method for the detection of a target substance in a sample. The probe array can be used for detecting a target substance with high reliability. Sequences AAC99241 - AAC99305 represent probes used in an array in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC99283 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18
                                                                                                                                                                                                                                                                                                                                  WPI; 2001-027424/04.
                                                                                                                                                                                                                                                                                                                                                                                28-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP2000270896-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Probe; probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           example illustrating the invention
                                                                                                                                                      Sequence 18
                                                                                                                                                                            example illustrating the invention
                                                                                                                                                                                                                                                 Example
                                                                                                                                                                                                                                                                        substance in a sample.
                                                                                                                                                                                                                                                                                                                                                                                                       28-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence
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                                                           18
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                                                                                                       12;
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                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATGAACCGGAGG 27
                                                           ATGAACCGGAGG 7
                                                                                ATGAACCGGAGG 27
                                                                                                                                                                                                                                                 Page 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                       Conservative
                                                                                                                                                    BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  array;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probe-combined
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in probe array
                                                                                                                                                                                                                                               20pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                  2.0%;
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Pred. No.
                                                                                                                  Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   substrate; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4; Le
. 1.2e+05;
                                                                                                                  DB 4; L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 18;
                                                                                                                            Length 18;
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                                                                                                       Indels
                                                                                                                                                                                                                                                                                              y, detection a single-
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                                                                                                      0
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X A X

AAC91122 AAC91122

standard;

DNA;

18

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WO200233068-A1

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                                                                                                                                                                                  RESULT 193
                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                         acid of a fungal pathogen possibly present in the sample with at least one oligonucleotide probe, from an Internal Transcribed Spacer (ITS) region. The method is useful for simultaneous detection and differentiation of clinically important fungi in a single assay, particularly Candida albicans, C. parapsilosis, C. tropicalis, C. kefyr, C. krusei, C. glabrata, C. dubliniensis, Aspergillus flavus, A. versicole, A. nidulans, A. fumigatus, C. neoformans and pneumocystis carinii. The method is especially useful in the detection of opportunistic infections in patients with impaired immunity systems, as organ transplants patients, patients receiving intensive anticancer treatments, diabetics or AIDS patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting and identifying fungal pathogens, especially Candida, Cryptococcus and Aspergillus, comprises hybridizing the amplified nucleic acid of the fungal pathogen with a probe from the internal transcribed spacer region of a DNA.
                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 3 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-061555/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-MAY-1999;
11-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fungal pathogenic; Internal Transcribed Spacer; opportunistic infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fungal
                                                                                                            13-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 46; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Smith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-MAY-2000; 2000WO-EP004714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200073499-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-MAR-2001
                                                           Nucleic acid
                                                                                                                                   ABK72521;
                                                                                   DNA sequence
                                                                                                                                                           ABK72521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (INNO-) INNOGENETICS NV.
                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                               510
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                                                                                                                                                                                                                                                                      l Similarity
12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pathogenic species identification probe
                                                                                                                                                           standard;
                                                                                                                                                                                                                                             regrererecae 521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maher M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               species in a sample. T
                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                    #19
                                                              base
                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99EP-00870109,
99US-0138621P.
                                                                                   relating to
                                                            sequence analysis; DNA diagnosis;
                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                 2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Martin C,
                                                                                                                                                            18
                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                      <u>.</u>.
                                                                                                                                                                                                                                                                                              Score 12;
                                                                                   nucleic
                                                                                                                                                                                                                                                                                   Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jannes G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The method involves hybridizing
                                                                                                                                                                                                                                                                       Mismatches
                                                                                    acid base sequence analysis method
                                                                                                                                                                                                                                                                                              DB 4;
                                                                                                                                                                                                                                                                                 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rossau
                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                              Length 18;
                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Van Der Weide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fungal
                                                                                                                                                                                                                                                                      <u>,,</u>
                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a nucleic
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Best Local S
Matches 12
           WPI; 2002-372310/40
                                                                                                                                                                               WO200233068-A1
                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a method of analysing an unknown nucleic acid base sequence. The method comprises preparing a probe array, hybridising with the probe array, measuring the fluorescence yield in the reaction, obtaining a template pattern, producing a sample pattern, and comparing the sample pattern with the template pattern. The method is useful for specifying an unknown base sequence at a defined site of a target single-stranded nucleic acid, which is useful for analysing a nucleic acid base sequence. The method is applicable in DNA diagnosis and therapy, and is useful in medicine and biology. Measuring the fluorescence yield allows the detection of a one-base mismatch which can be considered to produce high detection of a one-base mismatch which can be considered to produce high detection in thermostability is less important, and the judgement on each spot can be reliably carried out.
                                      Yamamoto N,
                                                                                                                                                      25-APR-2002
                                                                                                                                                                                                                                    Nucleic acid base sequence analysis; DNA diagnosis;
                                                                                                                                                                                                                                                                   Sample origonucleotide #42
                                                                                                                                                                                                                                                                                                                          ABK72480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18
                                                                                            18-OCT-2000;
                                                                                                                       18-OCT-2000; 2000WO-JP007244.
                                                                                                                                                                                                                                                                                                13-AUG-2002
                                                                                                                                                                                                                                                                                                                                                        ABK72480 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Screening an unknown base sequence at a defined site of a target single-
stranded nucleic acid for use in DNA diagnosis and therapy, comprises a
DNA chip, fluorescence yield and pattern-based method.
                                                                  (CANO ) CANON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK72503-ABK72524 represent DNA sequences described in the specification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-372310/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-OCT-2000; 2000WO-JP007244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-OCT-2000; 2000WO-JP007244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      16 ATGAACCGGAGG 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                            ATGAACCGGAGG 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 5 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                      Okamoto
                                                                                              2000WO-JP007244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 8; 53pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Okamoto
                                                                                                                                                                                                                                                                                              (first entry)
                                                                  츳
                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.0%;
                                      Suzuki
                                                                                                                                                                                                                                                                                                                                                        명
                                                                                                                                                                                                                                                                for analysing nucleic acid base sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 6; Le 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 18;
                                                                                                                                                                                                                                    probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Best
                                              Query Match
                                                                                                                                                                      nucleic acid base sequence. The method is applicable in DNA diagnosis and therapy, and is useful in medicine and biology. Measuring the fluorescence yield allows the detection of a one-base mismatch which can be considered to produce high detection accuracy. The hybrid pattern of the DNA probe is used so the difference in thermostability is less important, and the judgement on each spot can be reliably carried out.
                                                                                                                                                                                                                                                                                                            The present invention relates to a method of analysing an unknown nucl acid base sequence. The method comprises preparing a probe array, hybridising with the probe array, measuring the fluorescence yield in reaction, obtaining a template pattern, producing a sample pattern, an comparing the sample pattern with the template pattern. The method is useful for specifying an unknown base sequence at a defined site of a target single-stranded nucleic acid, which is useful for analysing a
                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Screening an unknown base sequence at a defined site of a target single-stranded nucleic acid for use in DNA diagnosis and therapy, comprises a DNA chip, fluorescence yield and pattern-based method.
                                                                                                                                                        important, and the ABK72439-ABK72502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 13; 53pp; Japanese.
                         Local
                       Similarity
                                                                                          18
                                                                                       BP; 2 A; 5 C; 6
                                                                                                                                                        represent sample
                    2.0%;
                                                                                       G; 5 T; 0 U; 0 Other;
                       Score 12;
Pred. No.
                                                                                                                                                      origonucleotides used in the present
                    DB 6; L
                                         Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     an unknown nucleic
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Matches
                                18
                                           16 ATGAACCGGAGG 27
                                                        12;
                                ATGAACCGGAGG
                                                         Conservative
                                7
ВP
                                                        0;
                                                        Mismatches
                                                      0;
                                                        Indels
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                                                       Gaps
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RESULT 195
ABK72522/c
ID ABK725
XX
AC ABK725
XX
DT 13-AUC
DT 13-AUC
XX
DT Nucle
XX
KW Nucle
XX
KW Nucle
XX
EN WO20(
XX
PN WO20(
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PN WO20(
XX
PD 25-AI
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PD Scr
PF 18-C
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PR WPI
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PI Yami
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PI Scr
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PT Scr
PT DN#
XX
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PT DN#
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CC Th
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II
                                                                                                                                                                                                                                                                                                                                 Nucleic acid base sequence analysis; DNA diagnosis; ds
                                                                                                                                                                                                                                                                                                                                                                                13-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                           ABK72522 standard; DNA; 18
                                                                                                                                                                                                                                                                                       WO200233068-A1
                                                                                                                                                                                                                                                                                                                                                       DNA sequence #20 relating to nucleic acid base sequence analysis method
                                                                                                                                                                                                                                                                                                                                                                                                       ABK72522
                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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reaction, hybridising

base sequence.

The present invention relates to a method of analysing an unknown nucleic

sequence. The method comprises preparing a probe array, my with the probe array, measuring the fluorescence yield in the obtaining a template pattern, producing a sample pattern, and the sample pattern with the template pattern. The method is r specifying an unknown base sequence at a defined site of a

Disclosure; Page 8; 53pp; Japanese

DNA chip,

Yamamoto N,

Okamoto

'n

Suzuki T;

(CANO) CANON KK

WPI; 2002-372310/40.

18-OCT-2000; 2000WO-JP007244. 18-OCT-2000; 2000WO-JP007244. 25-APR-2002

Screening an unknown base sequence at a defined site of a target single-stranded nucleic acid for use in DNA diagnosis and therapy, comprises a

fluorescence yield and pattern-based method.

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RESULT 196
ABN99788/c
ID ABN997
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                                                 Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                target single-stranded nucleic acid, which is useful for analysing a nucleic acid base sequence. The method is applicable in DNA diagnosis and therapy, and is useful in medicine and biology. Measuring the fluorescence yield allows the detection of a one-base mismatch which can be considered to produce high detection accuracy. The hybrid pattern of the DNA probe is used so the difference in thermostability is less important, and the judgement on each spot can be reliably carried out. ABK72503-ABK72524 represent DNA sequences described in the specification of the present invention
                                                                                                                                     The invention relates to a novel method for detecting a complex formed between a probe and its complement. The method is used for detecting a complex formed between an oligonucleotide of known basequence accomplementary probe, and for evaluating if the sequence is contained in liquid samples, or the level of binding by using the oligonucleotide as the detecting probe. The sequence represents a probe used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18
                                                                                                    Sequence
                                                                                                                                                                                                                               Example 3; Page 19; 25pp; Japanese.
                                                                                                                                                                                                                                                                      Detection of an object component in a sample using an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                  31-AUG-2000; 2000JP-00263395
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA probe #42
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                                                                                                                             Invention
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12; Conserv
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ATGAACCGGAGG 7
                         ATGAACCGGAGG
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                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                array; oligonucleotide detection; ss.
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                                                                                                    2 A;
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Pred. No. 1.2e+05;
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Pred. No.
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                                                    Mismatches
                                                                                                    T; 0 U; 0 Other;
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                                                                        Length 18;
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RESULT 197 ABL54942/c

/note= "Rhodamine labelled"

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RESULT 198
ABL54965
AX FX BX BX SX B F F F
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; p53; probe; variation detection; DNA array; ss.
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                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Screening for gene variation by using DNA array in which probes giving strong signals forming hybrids with normal sequence, and probes having sequences expected to form hybrids with variants are separately arrang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-AUG-2000; 2000JP-00263396
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                                                                                Human;
                                                                                                         Human rhodamine labelled p53 gene fragment #65 (normal sequence)
                                                                                                                                 18-JUN-2002
                                                                                                                                                         ABL54965
                                                                                                                                                                                ABL54965 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human tumour suppressor gene
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                                                          sapiens
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                                                                                 p53;
                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                             ATGAACCGGAGG 7
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                                                                                                                                                                                                                                                                                                                                          BP;
                                                                                 rhodamine; variation detection; DNA array;
                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                          2 A;
                                  Location/Qualifiers
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1.2e+05;
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RESULT 199
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequence represents a section of the normal p53 gene, designated DNA #65. The sequence was used as a target for hybridisation in the invention, and hybridises to probe #42. The invention relates to a novel method for screening for a variation in a nucleic acid sequence. The method involves using a DNA array in which a group of probes which will give strong signals forming hybrids with a normal gene sequence, and a group of probes having sequences expected to form hybrids with gene variants are separately arranged. The method is useful for screening for the presence or absence of variation in a nucleic acid sequence. The method is also useful for mass screening to determine rapidly the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Screening for gene variation by using DNA array in which probes giving strong signals forming hybrids with normal sequence, and probes having sequences expected to form hybrids with variants are separately arranged
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Base oligonucleotide for preparation of a probe.
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19-FEB-2001; 2001JP-00042344
                                                                                                                                                                                                                                              18-SEP-2001; 2001EP-00307932
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Liquid discharge; nucleic acid analysis; gene examination; probe; ss
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Pred. No.
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hes 0;
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                                      The invention relates to producing a probe support. The method involves (a) providing a liquid discharging device including reservoirs for containing liquids containing the probes and discharge mozzles connecting with the corresponding reservoirs; (b) aligning the discharge mozzles and the support relatively; and (c) discharging the liquids containing the probes from the discharge mozzles to different positions on the support. The number of reservoirs and discharge nozzles are the number of probes. The method is useful for producing probe supports useful in base sequence analysis of gene decoxyribonucleic acids (DNAs) and gene examination. The present sequence represents a base oligonuclectide used for preparation of a probe
Sequence
                                                                                                                                                                                                                                                                                                                                                                                              Producing probe supports for use in base sequence analysis of gene deoxyribonucleic acid, involves providing liquid discharging device for two-dimensionally arranging and fixing probe arrays on solid-phase
                                                                                                                                                                                                                                                                                                                             Disclosure; Page 20; 53pp; English
18
BP;
5 A;
6 C;
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<u>ი</u>
2 T; 0 U;
0 Other;
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                                        Query Match
Best Local S
                                  Matches
               16 ATGAACCGGAGG
\vdash
                               l Similarity
12; Conserv
ATGAACCGGAGG 12
                                 Conservative
                                         100.0%;
               27
                                                 2.0%;
                                 0;
                                         Score 12;
Pred. No.
                                 Mismatches
                                        DB 6; Le
1.2e+05;
                                                Length 18
                                 Indels
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                               Gaps
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RESULT 200
ABK88543
ID ABK885
XX ABK885
XX Probe
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12-DEC-2000; 2000JP-00377349.
                                          12-DEC-2000; 2000JP-00377349
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(CANO ) CANON KK
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Detecting nucleic acid with hybrid formation of a probe nucleic acid and a target nucleic acid without interference of the other double stranded nucleic molecules.

WPI; 2002-569947/61.

Example 1; Page 9; 13pp; Japanese

The invention describes a hybrid of a probe nucleic acid and a target nucleic acid for detection of nucleic acid. This sequence represents sythetic hybridisation probe used in the nucleic acid detection methors. invention a target method õ

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Query Match
Best Local Similarity
                            Sequence 18
                           BP; 5
                          A; 6 C; 5 G; 2 T; 0 U; 0 Other;
2.0%;
Score Pred.
12;
No.
DB 6; I
        Length
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18

WPI; 2002-364388/40

Yamamoto

Watanabe

ä

Suzuki

H,

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RESULT 202
ABT04735/c
ID ABT047
XX
ABT047735/c
AC ABT047
XX
DT 27-SEP
XX
End-la
XX
End-la
XX
End-la
XX
Uniden
XX
JP2002
XX
PD 28-MAY
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ABK88542/c
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Best Local S
Matches 12
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                                                                                                                                                                                                                                                                                                          The invention describes a hybrid of a probe nucleic acid and a target nucleic acid for detection of nucleic acid. This sequence represents a sythetic target gene fragment for probe hybridisation used in the nucleic acid detection method of the invention
                                                                                                                                ABT04735
                                                                                                                                                                                                                                                                                         Sequence 18
                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 9; 13pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                     nucleic molecules.
                                                                                                                                                                                                                                                                                                                                                                                             Detecting nucleic acid with hybrid formation of a probe nucleic acid and a target nucleic acid without interference of the other double stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK88542 standard;
      28-MAY-2002
                                             Unidentified
                                                                End-labelled probe array production; probe; ss; target substance
                                                                                       End-labelled
                                                                                                          27-SEP-2002
                                                                                                                                                   ABT04735 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-569947/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-DEC-2000; 2000JP-00377349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-DEC-2000; 2000JP-00377349
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP2002176999-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Probe nucleic acid; target nucleic acid; hybrid; nucleic acid detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic target gene fragment for nucleic acid detection method
                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CANO ) CANON
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                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                          ATGAACCGGAGG 27
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                                                                                                                                                                                                       ATGAACCGGAGG
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                                                                                                                                                                                                                                                                                        BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                      probe
                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                    DNA; 18
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                                                                                      array
                                                                                                          entry)
                                                                                                                                                                                                                                                        2.0%;
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                                                                                   production method-related oligonucleotide
                                                                                                                                                    48
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                                                                                                                                                                                                                                                        Score 12;
Pred. No.
                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                  DB 6;
                                                                                                                                                                                                                                                         1.2e+05;
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                                                                                                                                                                                                                                                                  Length 18
                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                     the nucleic
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                                                                 capture
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  RESULT 20
ABTIO4758
ID ABTIO
XX ABTIO
XX ABTIO
XX ABTIO
XX End-
XX End-
XX Unit
XX Unit
XX UP2:
PN JP2:
XX 24-:
PF 24-:
PF 24-:
XX Pre
PT Sub
XX Pre
PT Sub
CC Is
CC Sub
CC Syn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises a method for the synthesis of an end-labelled probe array - in which part of a probe for capturing a target substance is fixed at a plural of the matrix sites on the surface of a probe array substrate. In the method of the invention the units for constituting the probe are combined successively and, at the final stage of the successive synthesis, a labelling substance is combined to the end of the probe and extended to a desired chain length. The method of the invention is useful
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Matches
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Best Local
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                                                                                                                    WPI; 2002-552742/59
                                                                                                                                                                                                            24-NOV-2000; 2000JP-00357446
                                                                                                                                                                                                                                                                                                       28-MAY-2002.
                                                                                                                                                                                                                                                                                                                                                    JP2002153284-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                     End-labelled probe array production; probe; ss; target substance capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          End-labelled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABT04758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABT04758 standard; DNA; 18
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Example 1; Fig 1; 25pp; Japanese
                                                                  Preparation of an end-labelled probe array, for capturing a target
                                                                                                                                                                                                                                                          24-NOV-2000; 2000JP-00357446
                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Preparation of an end-labelled probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-552742/59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-NOV-2000; 2000JP-00357446.
                                                                                                                                                                 (CANO)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 2 A; 5 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       array
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       production method-related oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Japanese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12;
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1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for capturing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a target
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RESULT 204
ABQ81304/c
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                                        provides for the use of antagonists of P450 enzymes for the prevention or alleviation of a tachyphylactic response to administration of a vitamin D canalogue, corticosteroid or macrolactam to a patient, e.g. for the treatment of psoriasis. The underlying cause of tachyphylaxis was shown to be degradation of a drug in the patient, rather than desensitization or receptor down-regulation. Exposure of a patient to the drug for extended periods results in an increase in the expression of enzymes which are capable of metabolizing the drug. A method for treatment of tachyphylaxis therefore involves inhibiting the induced metabolic enzyme, especially a p450 cytochrome, by administration of an antagonist of the enzyme. Detection of an increase in the amount and/or activity of a metabolic enzyme capable of metabolizing a drug following extended exposure of a cell from an individual to the drug indicates the increased likelihood of that individual developing a tachyphylactic response to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                        Alleviating or preventing a tachyphylactic response to an agent and treating psoriasis, comprises administering an antagonist of a metabolic enzyme, which is induced as a result of exposure to the agent, to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-713234/77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-DEC-2000; 2000GB-00029524.
                                                                                                                                                                                                                                                                             analogues,
                                                                                                                                                                                                                                                                                                                                            Example 1; Page 75; 136pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-2001; 2001WO-GB005369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cytochrome P450; CYP2A6; enzyme; tachyphylaxis; drug tolerance; human; psoriasis; antipsoriatic; antipruritic; dermatological; PCR; primer; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for the production an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200245704-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytochrome P450 CYP2A6 sense primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABQ81304 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MOLE-) MOLECULAR SKINCARE LTD.
                                                                                                                                                                                                                                                                        present sequence is a sense primer for cytochrome P450 CYP2A6. RT-PCI used to characterise metabolic enzyme induction by vitamin D logues, corticosteroids and macrolactams in human skin. The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16 ATGAACCGGAGG 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bavik C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of a probe array.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.0%; Score 12;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cork M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tazi-Ahnini
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.2e+05;
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Query Match Best Local Similarity

2.0%;

Pred. No.

Length 18;

Sequence 18

BP; 8

A,

N C; 8

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년; 0

U; 0 Other; DB 6; Le

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RESULT 206
ABT06256/c
ID ABT062
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AC ABT062
XX
AC ABT062
XX
DT 24-OCT
XX
DE Synthe
XX
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ABL59677/c
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Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                   The present invention describes a method for determining simultaneously the reactivity of a first sample with other samples, in which the second to the 2 plus nth (n is not less than 1) samples having different properties are arranged independently on a substrate, on whose surface the first sample is already present, and the reactivities between the first sample and each of the second to the 2 plus n-th samples are determined. Also described is a tissue sample matrix in which several samples from different sources are present on each matrix divided on a substrate. The method is used for determining simultaneously the reactivity of a first sample with several other differing samples. ABL5965 to ABL59701 represent oligonucleotide probes used in an example from the present invention
            Synthetic DNA selling system -
                                                                ABT06256
                                       24-OCT-2002
                                                                                       ABT06256 standard; DNA; 18 BP
                                                                                                                                                                                                                                                               Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 11; 24pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Simultaneous testing of the reactivity of a sample with other different samples, comprises applying to the two samples to a substrate comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-AUG-2000; 2000JP-00263505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Simultaneous determination; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide probe SEQ ID NO:42.
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                                                                                                                                                          8
                                                                                                                                                                                 16 АТGААСССБВАСС 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16
                                                                                                                                                                                                          12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        matrices.
                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TIGCICTICCIC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TIGCICITCCIC 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                          2.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                              BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                               Þ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                            5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВÞ
                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                                                                      Score 12;
Pred. No.
            related oligonucleotide
                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                     DB 6; Le
                                                                                                                                                                                                                                  Length 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ..
                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                            RESULT 207
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention comprises a synthetic DNA selling system using the internet. The system displays a purchase order menu display, with the number of base sequences of DNA from which the orderer selects a DNA. The order information is transmitted to a successful bidder side server which orders for production and delivery of selected synthetic DNA. The system of the invention is useful for marketing synthetic DNAs of different base sequences and concentrations according to the desire of the user, especially genes concerned with human major histocompatibility complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          synthetic DNA selling system; internet; ss; major histocompatibility complex; MHC.
                      Grubman MJ,
                                                                     26-APR-2001;
24-APR-2002;
                                                                                                       26-APR-2002; 2002WO-US013247
                                                                                                                                  07-NOV-2002
                                                                                                                                                          WO200287336-A1
                                                                                                                                                                                                         Porcine; interferon; IFN; foot FMD; virucide; PCR; primer; ss.
                                                                                                                                                                                                                                             Porcine interferon
                                                                                                                                                                                                                                                                     02-MAY-2003
                                                                                                                                                                                                                                                                                             AAD51874;
                                                                                                                                                                                                                                                                                                                    AAD51874 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 5; 22pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the successful bidder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic DNA selling system using the Internet, displays purchase order menu to orderer's terminal and initiates production of selected DNA for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic DNA selling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-492955/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-AUG-2000; 2000JP-00259715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-AUG-2000; 2000JP-00259715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JP2002074089-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
2003-140146/13
                                                                                                                                                                                                                                                                                                                                                                                 18
                                                                                                                                                                                                                                                                                                                                                                                                        16
                                                                                                                                                                                                                                                                                                                                                                                                                                12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotides ABT06196 -
                                             Sn
                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                 ATGAACCGGAGG 7
                                                                                                                                                                                                                                                                                                                                                                                                      ATGAACCGGAGG 27
                                              SEC OF AGRIC
                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                     2001US-0286345P
2002US-00128463
                     Chinsangara
                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                          (IFN) -alpha DNA amplifying forward PCR primer
                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                          2.0%;
                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                   18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Japanese.
                                                                                                                                                                                                                                                                                                                   股
                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 12;
Pred. No.
                      Koster M,
                                                                                                                                                                                                                    and
                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                   mouth disease; vaccine; immunisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABT06278 are used in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6; Lt
                      Moraes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     purchase order menu.
                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 18;
                      Ŋ,
                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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ATGAACCGGAGG

Matches Query Match Best Local

l Similarity 12; Conserv

Conservative

0,

Mismatches

Indels

0;

Gaps

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100.0%;

Score 12; Pred. No.

DB 7; Lt

Length 18

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RESULT 208
ABZ21481
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local :
                         The invention relates to preparation of a probe array by high speed accourate dropping of the probe solution to improve quality of the parray. The probe array is useful in the analysis of base sequences and reliable genetic screening of multiple items. The present seque that of a probe used in examples of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mouth disease (FMD). The construct, vaccine and method are useful for protective immunisation against foot and mouth disease in animals. The construct is useful for preparity avaccine, which is particularly useful for stimulating protective response to such disease. The present sequence is porcine interferon (pIFN)-alpha DNA amplifying PCR primer. This sequence is used in the exemplification of the invention
                                                                                                                                       A process for preparation of a high density array of probes, used for DNA analysis and screening, comprising solution dropped on a carrier to form multiple spots at high speed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 4 A; 9 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel construct comprising a DNA sequence capable of expressing interferon (IFN) in animals susceptible to foot and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disease
Sequence
                                                                                                                                                                                                                                                                                                                   10-SEP-2002.
                                                                                                                                                                                                                                                                                                                                               JP2002253251-A
                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic probe SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ21481;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ21481 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New construct comprising a DNA sequence capable of expressing interferon in animals susceptible to foot and mouth disease, useful for preparing a
                                                                                                             Example 3; Page 12; 19pp; Japanese.
                                                                                                                                                                                                  WPI; 2003-096532/09.
                                                                                                                                                                                                                                                          28-FEB-2001; 2001JP-00055972
                                                                                                                                                                                                                                                                                     28-FEB-2001; 2001JP-00055972
                                                                                                                                                                                                                                                                                                                                                                                                     Probe array; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3;
                                                                                                                                                                                                                              (CANO ) CANON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ຫ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for providing protective immunization against
 <u>1</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCCCAACCTCAG 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CCCCAACCTCAG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 71;
 B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                     88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               72pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.0%;
6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ٠,
د
                                          sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                   probe
                                            DNA
is
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RESULT 210
ABV77868
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ABZ21482/c
ID ABZ214
SSXSXBXBX8X8
                                                                                                                                                                                                                                                       맑
                                                                                                                                                                                        Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                맑
                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                             analysis and screening, compr
multiple spots at high speed.
                                                Oligonucleotide SEQ ID 1 used to prepare a
                                                                                                              ABV77868 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                    A process for preparation of a high density array of probes, used for DNA analysis and screening, comprising solution dropped on a carrier to form
                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-096532/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-FEB-2001; 2001JP-00055972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-SEP-2002
                            Probe; probe carrier; DNA micro-chip;
                                                                                            ABV77868;
                                                                                                                                                                                                                                                       Sequence 18
                                                                                                                                                                                                                                                                                 The invention relates to preparation of a probe array by high speed and accurate dropping of the probe solution to improve quality of the probe array. The probe array is useful in the analysis of base sequences of DNA and reliable genetic screening of multiple items. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP2002253251-A
                                                                       24-FEB-2003
                                                                                                                                                                                                                                                                                                                                       Example 3; Page 12; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2001; 2001JP-00055972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Probe array; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ21482;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ21482 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                            (CANO ) CANON KK
                                                                                                                                                                                                                                                                          of a probe used in examples of the invention
                                                                                                                                                                   18 ATGAACCGGAGG 7
                                                                                                                                                                                        16
                                                                                                                                                                                                             12;
                                                                                                                                                                                        ATGAACCGGAGG 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probe SEQ
                                                                                                                                                                                                           2.0%; So ilarity 100.0%; I Conservative 0;
                                                                                                                                                                                                                                                      BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/mod_base= OTHER
/note= "G-OPO3-(CH2)6-SH"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N
                                                                                                                BP.
                                                                                                                                                                                                                       Score 12;
Pred. No.
                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                           1.2e+05;
hes 0;
                                                                                                                                                                                                                                DB 7;
                                                 probe
                                                                                                                                                                                                                                Length 18;
                                                                                                                                                                                                             Indels
                                                                                                                                                                                                           0
                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                     RESULT 211
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                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a device for manufacturing a probe carrier with some types of probes. The device comprises: (i) a solution discharge unit composed of reservoirs for housing the respective types of probe solutions; (ii) groups of discharging openings connected to the respective reservoirs; and (iii) discharging energy generating members corresponding to the respective discharging openings one by one. The device is suitable for manufacturing probe carriers, such as, DNA microchips etc. used for analysing base sequence of genetic DNA etc. The present sequence is a oligonucleotide used for preparing a probe, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Device for manufacturing a probe carrier used for analyzing of genetic DNA comprises discharging liquid through a group discharging openings on each probe spot.
                      28-FEB-2001; 2001JP-00055970.
                                              11-SEP-2002
                                                                       JP2002257694-A
                                                                                                                                                                       Synthetic.
                                                                                                                                                                                            Probe; probe carrier; DNA micro-chip; ss
                                                                                                                                                                                                                     Oligonuclectide SEQ ID 2 used to prepare a probe.
                                                                                                                                                                                                                                              24-FEB-2003
                                                                                                                                                                                                                                                                      ABV77869
                                                                                                                                                                                                                                                                                             ABV77869 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 5 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-079066/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; Page 12; 18pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP2002257694-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-FEB-2001; 2001JP-00055970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CANO)
                                                                                                                                                                                                                                                                                                                                                                                  16
                                                                                                                                                                                                                                                                                                                                                                                                          12;
                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                          ATGAACCGGAGG
                                                                                                                                                                                                                                                                                                                                                                                  ATGAACCGGAGG
                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                              (first
                                                                                                                                           Location/Qualifiers
                                                                                             /mod_base=
/note= "5'
                                                                                                                       /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "5' labelled with tetramethyl rhodamine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                   2.0%;
                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                          12
                                                                                                                                                                                                                                                                                                                                                                                  27
                                                                                             : OTHER
labelled
                                                                                                                                                                                                                                                                                             ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                         °,
                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                               with
                                                                                                                                                                                                                                                                                                                                                                                                                    DB 7; L
                                                                                             SH- (CH2) 6-PO4 "
                                                                                                                                                                                                                                                                                                                                                                                                                               Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       base sequence of
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28-FEB-2001; 2001JP-00055970

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chips etc. used for analysing base sequence of genetic DNA etc. The present sequence is a oligonucleotide used for preparing a probe, which was used in an example from the invention. Note: This sequence is shown in this orientation (5' to 3') in the disclosure of the specification, but is shown in the opposite orientation in the sequence listing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the present inventor to a solution carrier with some types of probes. The device comprises: (i) a solution discharge unit composed of reservoirs for housing the respective types of probe solutions; (ii) groups of discharging openings connected to the respective reservoirs; and (iii) discharging energy generating members corresponding to the respective discharging openings one by one. The device is suitable for manufacturing probe carriers, such as, DNA microdevice is suitable for manufacturing probe carriers, such as, DNA microdevice is suitable for manufacturing probe carriers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Device for manufacturing a probe carrier used for analyzing of genetic DNA comprises discharging liquid through a group discharging openings on each probe spot.
The invention comprises a method for manufacturing a probe carrier. The method involves discharging and fixing probe solutions in the corresponding cavities according to information on the positions of respective types of probes. The method of the invention is useful for manufacturing probe carriers (e.g. DNA micro-chips). The present DNA
                                                                                                                                                    Manufacturing a probe carrier used for analyzing a base sequence of genetic DNA, comprises discharging a probe solution in a cavity located at each probe position on a probe carrier through a group of discharging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Probe carrier manufacturing method-related oligonucleotide, SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-MAR-2003
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                                                                                                        Example 4;
                                                                                                                                                                                                                           WPI; 2003-106341/10
                                                                                                                                                                                                                                                                                          28-FEB-2001; 2001JP-00055971
                                                                                                                                                                                                                                                                                                                           28-FEB-2001; 2001JP-00055971
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                                                                                                      Page 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
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RESULT 214
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                                                                                                                                                                                                            The invention relates to a novel probe carrier and the method for manufacturing the carrier. The invention enables stable discharge of solution, and removes liquid droplets adhering to discharge nozzle. present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                          Probe carrier manufacturing method for inkjet liquid discharge head in direction orthogonal angle satisfying predetermined relation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence represents an oligonucleotide invention
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                                   18-DEC-2003
                                                     ADC54007;
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                                                                                                                                                                                                                                                          Example 2;
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. 1.2e+05;
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AAT43666

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Best Local S
Matches 12
                                                                                             Human immunodeficiency virus antibodies - useful for asses
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Probe carrier manufacturing method for inkjet system, involves scanning liquid discharge head in direction orthogonal to scanning direction, at angle satisfying predetermined relation.
                                                                                                                                                                              WPI; 1996-507733/51
                                                                                                                                                                                                                                                                                                                                    18-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                matrix protein p17; prognosis; prohybridisation assay; immunoassay;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1; subtype B; transmissible; matrix protein p17; prognosis; probe; detection; maternal transmission;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-1 matrix protein p17 gene
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Matches Query Match Best Local S

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BP; 1 A; 7 C; 2 G; 9 T; 0 U; 0 Other;

infection

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RESULT 216
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Best Local
                                                      A set of six oligonucleotide probes enable the risk of maternal- foetal transmission of subtype B HIV-1 to be evaluated. Probes 1 and 6 (see NAT43664 and AAT43669) allow detection of HIV-1 strains which are always transmitted to the foetus; probes 2-4 (see AAT43665-T43668) allow detection of strains which are never transmitted from mother to foetus. The present sequence is that of probe 2 and it detects HAR, LOUB, VIL and 2754 HIV-1 sequences
                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus p17 gene fragments, derived proteins and antibodies - useful for assessing the risk of maternal transmission of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-507733/51.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Narwa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (COMS ) COMMISSARIAT ENERGIE
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                                                                                                                                                                                                                                                                                                       15;
                                                                                                                                                                                                                                                                                                                                                                infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TIGCTCTTCCTC 186
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                                                                                                                                                                                                                                                                                             Page 29; 46pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Roques
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   assay; immunoassay; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95FR-00005914.
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P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ν.
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1.2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           maternal transmission;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>.</u>
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RESULT 218
AAT68898/c
ID AAT688
XX AAT688
XX ACAF688
XX O6-APR
XX Human
XX Drug-I
KW Drug-I
KW Droger
XX Dynthe
OS Homo :
XX PN WO971:
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AAT43667
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                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                      A set of six oligonucleotide probes enable the risk of maternal foetal transmission of subtype B HIV-1 to be evaluated. Probes 1 and 6 (see AAT43664 and AAT43669) allow detection of HIV-1 strains which are alway transmitted to the foetus; probes 2-4 (see AAT43665-T43668) allow detection of strains which are never transmitted from mother to foetus. The present sequence is that of probe 4 and it detects the CEL HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus p17 gene fragments, derived proteins and antibodies - useful for assessing the risk of maternal transmission of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1996-507733/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus type matrix protein p17; prognosis; pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV-1 matrix protein p17 gene probe 4.
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                                                                                               Human beta-actin 5'
                                                                                                                       06-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; Page 29; 46pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP743364-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hybridisation assay; immunoassay; ss.
                                                                                                                                               AAT68898
                                                                                                                                                                     AAT68898 standard;
                                                                                                                                                                                                                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-NOV-1996
                        Homo sapiens
                                                                        Drug-resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-MAY-1996;
                                                                                                                                                                                                                                                    175 Trecretrecte 186
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                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96EP-00401084
                                                          neoplastic disease; non-malignant haematopoietic cell;
rearrangement; RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                         A;
                                                                                                                                                                    DNA;
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                                                                                               RT-PCR
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G;
                                                                                              primer.
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                                                                                                                                                                                                                                                                                     Score 12; DB 2; Li
Pred. No. 1.2e+05;
                                                                                                                                                                                                                                                                                                                        9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probe;
                                                                                                                                                                                                                                                                            Mismatches

    subtype B; transmissible;
    detection; maternal tran

                                                                                                                                                                                                                                                                                                Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transmission;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        promoter which functions in the host cell. The second nucleic acid molecule operably linked to a second molecule encodes an RNA molecule or a polypeptide whose expression decreases the expression of an RNA or a polypeptide present in a malignant cell only. This method can eliminate residual neoplastic disease in a patient, where the disease has an immature hematopoietic progenitor cell with a well-defined gene rearrangement. Diseases such as chronic myelogenous leukaemia which is associated with a BCR/ABI gene rearrangement, acute lymphoblastic leukaemia and acute promyelocytic sequence 19 pp.
                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human beta-actin 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT68899;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        drug-resistant, non-malignant haematopoietic cells. This method involves the construction of a new expression cassette comprising a first nucleic acid molecule which encodes resistance of a host cell to a cytotoxic agent, operably linked to a first promoter which functions in the host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             decreases RNA or protein found only in malignant cel
leukaemia(s), such as chronic myelogenous leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Expression cassette for forming drug resistant hematopoietic stem cells decreases RNA or protein found only in malignant cells; for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-289281/26.
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                                                                                                           14-NOV-1995;
                                                                                                                                                                                                                    22-MAY-1997
                                                                                                                                                                                                                                                                         WO9718305-A2
                                                                                                                                                                                                                                                                                                                                                                                                           progenitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Drug-resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This beta-actin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Fig 1; 52pp; English.
                                                                                                                                                                13-NOV-1996;
                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MINU ) UNIV MINNESOTA.
                                                       (MINU ) UNIV MINNESOTA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGAGGATCTTCA 328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 6 A; 6 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                           gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                      neoplastic disease; non-malignant haematopoietic cell; rearrangement; RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RT-PCR primer is used in a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.08; **
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RT-PCR primer.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local |
                                        Detection of chromosomal abnormalities - by subjecting patient nucleic acids to a multiplex molecular amplification procedure primers specific for characteristic nucleic acid sequence.
                                                                                                                                                                                                          06-DEC-1996;
                                                                                                                                                                                                                                          08-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                PCR primer; chromosomal abnormality; abnormality detection; leukaemia; lymphoma; carcinoma; adenocarcinoma; sarcoma; glioma; neuroblastoma; medullablastoma; malignant melanoma; malignant neoplastic condition; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV40952
                                                                                                                                           Pallisgaard N,
                                                                                                                                                                                                                                                                                                        W09824928-A2
                                                                                                                                                                                                                                                                                                                                           Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer BCR1ABL:1698U19 for abnormality detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-SEP-1998
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12; Conserv
                                                                                                                                                                          PALLISGAARD N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGAGGATCTTCA 328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 6 A; 6 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                          96DK-00001401
                                                                                                                                                                                                                                          97WO-DK000556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 19;
                                                                                                                                                                                                                                                                                                                                                                                    condition; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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74; 126pp; English

DNA sequence coding for oncofetal ferritin 1 protein, useful for immunizations against breast cancer, for enhancing fertilization during in vitro fertilization treatment and for use as a growth

factor

2000-271427/23

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RESULT 221
AAA136
XX
AAA136
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AC AAA136
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BU 20-JUI
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Human
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Human
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Human
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For leuke
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For Spon'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC This sequence represents a primer used in the method of the invention for the detection of the presence or absence of chromosomal abnormalities, ce each abnormality being associated with a condition in a subject and each combined abnormality being associated with a condition in a subject and each come characteristic nucleic acid sequence. The come thouse comprises: (a) obtaining a sample of nucleic acids derived from a subject which may harbour one of the chromosomal abnormalities; (b) come the sample to a multiplex molecular amplification (MMA) compresedure, where a number of the characteristic sequences, if present in carefular amplification (MMA) compresedure, where a number of the characteristic sequences, if present in carefular amplification (MMA) and detecting the presence and/or absence of an amplicon compresent to the presence or capacity of the abnormal sequences to detect the presence or capacity of the abnormal sequences to detect the presence or capacity of the abnormal sequences to detect the presence or capacity of the proprises the use of at least 7 mutually distinct primers (MDP) in one single reaction mixture, each of the primers defining an end of at least one characteristic nucleic acid sequences, the characteristic nucleic acid sequences, the characteristic nucleic acid sequences ach being capacity of the MDP. The methods can be used for detecting chromosomal abnormalities associated with diseases including numerous leukaemia's, lymphoma's, carefulariant and malaroma's, gliona's, neuroblastoma's, carefulariant and malaroma's, gliona's, neuroblastoma's,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pathological pregnancy; breast cancer; cytostatic; immunos; contraceptive; abortive; nootropic; vaccine; immunisation; transplant rejection; autoimmune disease; fertilisation; d
                                                                                                                                                                                                                                                                                                                                                                                                11-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WC200015788-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in vitro fertilization; IVF; heptablastoma; Hodgkin's lymphoma; leukaemia; non-Hodgkin's lymphoma; embryonal tumour; Down's Syndrome; spontaneous abortion; miscarriage; premature contraction; toxaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human oncofoetal ferritin 1 PCR primer #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA13623 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 3 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                medullablastoma, malignant melanoma, and malignant neoplastic
                                                                                                                                                                                                                                                                                                                              (GARD-) GARDINO INVESTMENT NV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               290 TTCTGCGAGGGA 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17 TTCTGCGAGGGA 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oncofoetal ferritin 1; OFF1; ferritin; transplantation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        delivery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                98IL-00126181.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-IL000485.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2; L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunosuppressive
sation; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
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Claim 29;

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49; 66pp;

English.

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RESULT 222
AAZ76946/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a specifically claimed PCR primer used in the isolation of oncofetal ferritin 1 (OFF1) protein. OFF1 has cytostatic, immunosuppressive, contraceptive, abortive and nootropic activities, and can be used as a vaccine. Compositions comprising the expression vector containing an OFF1 coding sequence, and the OFF1 coding in the treatment of transplant rejections, autoimmune diseases, pathological pregnancies; (c) for enhancing fertilisation rates convert of bone-marrow progenitor cells such as granulocyte monocytes. The OFF1 nucleotide sequence is useful for diagnosing cancer such as breast cancer, heptablastoma, leukaemia, Hodgkin's and non-CC Hodgkin's lymphomas and embryonnal tumours, Down's Syndrome, and pathological pregnancies such as spontaneous abortion and miscarriage, comparature contractions, toxaemia or premature delivery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                     genomic map; haplotype; phenotype; polymorphic base; genotyping haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; identification; snp; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human genome;
genomic map; h
                                                                                                                                                                                                                                                21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human biallelic marker downstream amplification primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ76946 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19
                                                                                                                                                                                       Cohen
                                                                                                                                                                                                                                                                                          21-APR-1999;
                                                                                                                                                                                                                                                                                                                      28-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                   WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                   (GEST )
                                                                                                                of the human
                                                                                                                                                            2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       246
                                                                                                                               biallelic markers used
                                                                                                                                                                                       'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7 CTCCTGGAGCCC 18
                                                                                                                                                                                                                   GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CICCIGGAGCCC
                                                                                                                                                                                       Blumenfeld
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 3 A; 8 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  biallelic marker; high
                                                                                                                                                                                                                                                98US-0082614P
98US-0109732P
                                                                                                                                                                                                                                                                                          99WO-IB000822
                                                                                                               genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.0%;
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Pred. No.
                                                                                                                               construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 density disequilibrium map;
olymorphic base; genotyping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.2e+05
                                                                                                                               ρ
                                                                                                                              high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ID NO:11302.
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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the

Sequence 19

BP; 4

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6 T;

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diseases

Claim

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Page

2640;

2745pp;

English

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RESULT 223
AAC70391
ID AAC70391
XX AAC703
XX AAC703
XX Single
XX Single
XX Single
XX Single
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XX H
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Matches
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                                                                                      The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in hum genes. These SNPs can be used in disease diagnosis and prediction of individual's susceptibility to disease, in forensic and paternity tes and in genetic mapping. In particular, the SNPs of the invention can used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson
                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single nucleotide polymorphism; SNP; human; genetic disease disease susceptibility; cardiovascular system; endocrine uneurological system; forensic testing; paternity testing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                        Claim 8;
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Lipshutz RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single nucleotide polymorphism PCR primer #148.
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                                                                                                                                                                                                                                                                                                                                                                                                                 genetic analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-611722/58.
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12; Conserv
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                                                                                                                                                                                                                                                                                                                                                     Fig 5; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; DNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGCTCTTCCTCC 4
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Patil N, 9
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3klar P;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 3; Le
. 1.2e+05;
ches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 19;
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                                                                                                                                                                                                           paternity testing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer;
                                                                                                                                                                                                                                                                       in human
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RESULT 224
AAC70385
ID AAC703
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                                                                                        Query Match
Best Local S
Matches 12
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Best Local S
Matches 12
                                                                                                                                                The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                  Altshuler D,
Lipshutz RJ,
                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                         Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                                                       31-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                        30-MAR-2000; 2000WO-US008440
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR pr
                                                                                                                                                                                                                                                                                                 WPI; 2000-611722/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo
                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                            WO200058519-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single nucleotide polymorphism PCR primer #144.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC70385;
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                                                                       300
                                                                                                                                                                                                                                                                                                                                                                                                                                             sapiens
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                                                                                                                                                                                                                                    œ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GAACCCCAACCT 12
                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                            AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                     WHITEHEAD INST BIOMEDICAL RES
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                                                    GAACCCCAACCT 12
                                                                     GAACCCCAACCT 311
                                                                                                                                                                                                                                Fig 5; 214pp; English
                                                                                       ilarity 100.0%;
Conservative
                                                                                                                          BP; 4 A; 7 C;
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                                                                                                                                                                                                                                                                                                                  Cargill M,
Patil N, '
                                                                                                                                                                                                                                                                                                                                                                       99US-0127248P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
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                                                                                                2.0%; Score 12;
.00.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                  Sklar
                                                                                                                          2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                          Daley GQ,
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Pred. No.
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                                                                                        Mismatches
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                                                                                       1.2e+05;
hes 0;
                                                                                                        DB 3;
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1.2e+05;
                                                                                                                                                                                                                                                                                                                          JS,
                                                                                                       Length 19;
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                                                                                                                                                                                                                                                                                                                          Lander ES;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer;
                                                                                       Gaps
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AAD18285
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                                                                                                                                                                                                        Matches
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Best Local
                                                                                      18-DEC-2001
                                                                                                        AAD18285
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RESULT 22 AAC70388 ID AAC7 XX

225

AAC70388

standard;

DNA;

19

ВP

Homo sapiens

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The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testir and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                    Human; cytostatic; cancer-related antigen; homeobox protein; HOXA7; HOXB7; ADP ribosylation factor 1; Arf-1; ATP dependent iron transporter; ABC-7; neoplastic process; ovarian cancer; benign serous cystadenoma;
                                                                                                                          Human beta-actin amplifying PCR primer #2.
                                                                                                                                                                                                                                                                               AAD18285 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 4 A; 7 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer;
vaccine; beta-actin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 8; Fig 5; 214pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genetic analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-611722/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Altshuler D,
Lipshutz RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC70388;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200058519-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            300 GAACCCCAACCT 311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                             GAACCCCAACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                              (first
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Patil N, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0127248P
                                                                                                                                                                                                                                                                               DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.0%;
     PCR
                                                                                                                                                                                                                                                                                                                                                                                                          12
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  primer;
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klar P;
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Pred. No.
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မ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
     88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 3; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 19,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lander
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ntified in human prediction of an paternity testing
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RESULT 227
ABL58227
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel cancer-related antigen useful for prognosis, diagnosis and treatment of cancer, especially ovarian cancer in an individual, comprises a fragment isolated from bacteriophage lambda.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 3 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                  A. tumefaciens
                                                                                                                                                                                                                      15-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-MAR-2000; 2000US-0189226P.
28-DEC-2000; 2000US-0258452P.
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                   05-SEP-2001
                                                                                        Agrobacterium
                                                                                                                        virulence protein;
                                                                                                                                           VirE2; cytostatic;
                                                                                                                                                                                                                                                           ABL58227,
                                                                                                                                                                                                                                                                                             ABL58227 standard; DNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                          317 TGAGGATCTTCA 328
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                    (first entry)
                                                                                        tumefaciens
                                                                                                                                                                                VirE2 DNA specific oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human beta-actin DNA
                                                                                                                                          gene therapy; transfection; transformation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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ARBSULT 228
AAP848
ID AAP848
XX AAF848
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CT 09-JUL
XX PCR pr
XX Human;
XX O9-MAR
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XX O9-MAR
XX O3-SEP
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XX WPI;
ZX WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                allowing the molecule to cross the membrane through the channel. The method is useful in delivering materials across membranes, particularly into cells or organelles, and subsequently producing transiently or stably transfected or transformed cells. The method is especially useful for delivering nucleic acid to a cell to achieve expression of a desired transgene by the cell, and for treating cancer cells, relying on a new non-viral system without LTR and the possible hazards connected with them. The present sequence represents a VirE2 specific oligo used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Delivering a material across a membrane, useful for producing transiently or stably transfected or transformed cells, comprises introducing Agrobacterium VirE2 into a membrane, where VirE2 forms a channel through
                                                                                                                                                                                                                                                                                                             Human; SPG4 gene; spastin; PSF-AD; gene therapy; autosomal dominant familial spastic paraplegia; I
                                                                                                                                                                                                                                                                                                                                                                                                   09-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 4 A; 3 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        membrane. The method involves introducing Agrobacterium VirE2, its fragment or homologue into a membrane, where VirE2, its fragment or homologue forms a channel through the membrane; contacting the membrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-618844/72.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAR-2000; 2000GB-00005005
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                              WPI; 2001-283966/30.
                                                                                                                                      03-SEP-1999;
                                                                                                                                                                                                          09-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF84826 standard; cDNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          homologue forms a channel through the membrane; contacting the mem with a molecule desired to be transferred across the membrane; and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of delivering a material across a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 21; 30pp; English
                                                                  Weissenbach J,
                                                                                                                                                                        03-SEP-1999;
                                                                                                                                                                                                                                           FR2798138-A1
                                                                                                                                                                                                                                                                                                                                                                 PCR primer used for RACE-PCR reactions of human SPG4 cDNA
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New human nucleic acid from the SPG4 gene, useful e.g.

for diagnosis of

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ABK89543/c
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                                                                                                                                                                                                                                                                                                                                                                                                                           Protecting plant from barley yellow dwarf virus infection, by transforming modified nucleic acid expressing translationally-altered RNA into a plant cell, which confers resistance against virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Barley yellow dwarf virus infection; BYDV; plant resistance; monocotyledon; wheat; Triticum; Sorghum; rice; Oryza; barley; Hordeum; maize; Zea; rye; Secale; triticale; cat; Avena; antiviral; gene therapy;
    plant resistance against infection with BYDV. The method is useful protecting a plant from BYDV infection, where the plant is a monocotyledon selected from wheat (Triticum), sorghum (Sorghum), ri
                                                                                                            comprises transforming a modified nucleic acid molecule into a plant cell, where the expression of the nucleic acid molecule results in the expression of a translationally altered RNA molecule which confers to texpression of a translationally altered RNA molecule which confers to texpression of a translationally altered RNA molecule which confers to texpression of a translationally altered RNA molecule which confers to texpression of a translationally altered RNA molecule which confers to texpression of a translationally altered RNA molecule which confers to the confers
                                                                                                                                                                                                                         The present invention relates to a new method of protecting a barley yellow dwarf virus (BYDY) infection. The method of the
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21-MAY-2001; 2001US-0292778P
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Best Local
                                             The invention relates to double-stranded oligonucleotides consisting of two complementary strands (Ia; Ib), each having 1-5 unpaired nucleotides, at either at their 3' and 5' ends, forming single-stranded overhangs. One of (Ia) and (Ib) is complementary to a target sequence, (DNA or RNA), that is to be specifically repressed. The oligonucleotides are preferably double stranded inhibitor RNA molecules with a couple of thymidine bases attached at the 3' or 5' ends. The targets are preferably nucleic acids that, when repressed, induce apoptosis, necrosis or differentiation of tumour cells and/or inhibit division of such cells. Typical of many specified targets include: genes of the BCL2 family, genes that encode
metalloproteases (matrix or membrane) or their inhibitors; genes encoding mutant forms of nuclear hormone receptors; a sequence encoding the Hifl-alpha transcription factor; sequences encoding various isoforms of the
                                                                                                                                                                                                                                           Disclosure;
                                                                                                                                                                                                                                                                                         New double-stranded oligonucleotides, useful e.g. for treatment and diagnosis of tumors, comprise complementary strands with single-stra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     differentiation; tumour cell division; BCL2 gene family; matrix metalloprotease; membrane metalloprotease; nuclear hormone receptor; transcription factor; vascular endothelial growth factor; p53; chromosomal translocation; leukaemia; androgen receptor; diagnosis; cancer; human papilloma vi AIDS; BSE; CJD; macular degeneration; angiogenesis; diabetic retino
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10-APR-2002; 2002FR-00004474.
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Pred. No.
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1.2e+
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diabetic retinopathy;
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                                                                                                                                                                                                                                                                                         single-stranded
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RESULT 231
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nuclear hormone receptor; transcription factor; vascular endothelial growth factor; p53; chrosomal translocation; vascular endothelial growth factor; p53; chrosomal translocation; leukaemia; androgen receptor; diagnosis; cancer; human papilloma virus; AIDS; BSE; CJD; macular degeneration; angiogenesis; diabetic retinopathy; psoriasis; rheumatoid arthritis.
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10-APR-2002;
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WPI; 2003-441571/41.
                                                                                                           Harel-Bellan
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2002FR-00004474.
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                                                                                                           Ait-Si-Ali S,
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0; Mismatches
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diagnosis of tumors, comprise complementary strands with single-stranded
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Disclosure; Page 17; 148pp; French.

CC double stranded inhibitor RNA molecules with a couple of thymidine bases CC attached at the 3' or 5' ends. The targets are preferably nucleic acids CC that, when represed, induce apoptosis, necrosis or differentiation of CC thmour cells and/or inhibit division of such cells. Typical of many CC specified targets include: genes of the BCL2 family; genes that encode CC metalloproteases (matrix or membrane) or their inhibitors; genes encoding mutant forms of nuclear hormone receptors; a sequence encoding the Hifl-CC alpha transcription factor; sequences encoding various isoforms of the CC vascular endothelial growth factor; viral genes, genes that express a CC mutated protein, e.g. inactive p53; genes that are formed by a CC chromosomal translocation, e.g. where associated with leukaemia, and CC genes that express androgen receptors. The oligonucleotides are used: (i) conditions caused by expression of a harmful gene or fusion protein, c.g. specifically cancer (e.g. associated with expression of mutant p53 or of CC the human papilloma virus B6 protein), viral infections, especially AIDS CC reancer-inducing viruses, or unconventional infections, especially AIDS CC and (ii) for treating hypervascular diseases, e.g. age-related macular degeneration, angiogenesis in tumours, diabetic retinopathy, psoriasis CC and rheumatoid arthritis. They may also be used in vitro, e.g. for treating transplants and for establishing a genetic profile, for conditional caused by expression of setable than either hybrids prepared from DNA or single-stranded CC immunological or intolerance represents the sequence in the effectively than plandids. This sequence represents the sequence in the effectively than plasmids. This sequence represents the sequence in wild type human p53 gene around the position of mutation R248W. The oligonucleotides of the invention can be targeted to this sequence. at either at their 3' and 5' ends, forming single-stranded overhangs. One of (Ia) and (Ib) is complementary to a target sequence, (DNA or RNA), that is to be specifically repressed. The oligonucleotides are preferably The invention returned (Ia, Ib), each naving incle-stranded two complementary strands (Ia, Ib), each naving incle-stranded (D) The invention relates to double-stranded oligonucleotides consisting of two complementary strands (Ia; Ib), each having 1-5 unpaired nucleotides,

Sequence 19 BP; 5 ď 6 C; 6 G; 2 T; 0 U; 0 Other;

S Matches Query Match Local 16 l Similarity
12; Conserv ATGAACCGGAGG 27 Conservative 100.0%; 2.0%; 0, Score 12; DB 7; Pred. No. 1.2e+0 Mismatches 1.2e+05; <u>,</u> Length 19 Indels ٥, Gaps 0

RESULT 232 ADA03187 ADA03187

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ATGAACCGGAGG 14

standard; DNA; 19

06-NOV-2003 (first entry)

Wild type sequence in human p53 around position R248Q in mutant gene.

nuclear hormone receptor; transcription factor; vascular endothelial growth factor; p53; chromosomal translocation; vascular endothelial growth factor; p53; chromosomal translocation; leukaemia; androgen receptor; diagnosis; cancer; human papilloma virus; AIDS; BSE; CJD; macular degeneration; angiogenesis; diabetic retinopathy; differentiation; tumour cell division; BCL2 gene matrix metalloprotease; membrane metalloprotease; antidiabetic; antipsoriatic; antirheumatic; antiarthritic; inhibitor repression; inhibitor RNA; apoptosis; necrosis; differentiation; tumour cell division; BCL2 gene family; cytostatic; virucide; anti-HIV; neuroprotective; ophthalmological; rheumatoid

Homo sapiens

New double-stranded oligonucleotides, useful e.g.

for treatment and

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RESULT 233
ADB54361/c
ID ADB543
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                                                                                                                                                                                                                                                                   CC double stranded inhibitor RNA molecules with a couple of thymidine bases CC that, when repressed, induce apoptosis, necrosis or differentiation of the tumour cells and/or inhibit division of such cells. Typical of many CC specified targets include: genes of the BCL2 family; genes that encode metalloproteases (matrix or membrane) or their inhibitors; genes encoding mutant forms of nuclear hormone receptors; a sequence encoding the Hifl-CC alpha transcription factor; sequences encoding various isoforms of the mutant forms of nuclear hormone receptors; a sequence encoding the Hifl-CC vascular endothelial growth factor; viral genes; genes that express a cc mutated protein, e.g. inactive p53; genes that are formed by a chromosomal translocation, e.g. where associated with leukaemia, and CC genes that express androgen receptors. The oligonucleotides are used: (i) to study gene function; (ii) for therapy or diagnosis, particularly of CC conditions caused by expression of a harmful gene or fusion protein, specifically cancer (e.g. associated with expression of mutant p53 or of the human papilloma viruse, or unconventional infections, especially AIDS or cancer-inducing viruses, or unconventional infections, e.g. BSE or CDD; and (iii) for treating hypervascular diseases, e.g. age-related macular degeneration, angiogenesis in tumours, diabetic retinopathy, psoriasis can rhammatoid arthritis. They may also be used in vitro, e.g. for conditionalization, or modification, of treatment regimes. They provide more stable than either hybrids prepared from DNA or single-stranded common pagenesis of the invention and they enter tumour cells more common conditions of the provide macune represents the sequence in the colliponucleotides of the invention can be targeted to this sequence.
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                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                   Sequence 19
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10-APR-2002;
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                                                                                                                                                                                                          Similarity
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                                                                                                             ATGAACCGGAGG 14
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                                                                                                                                                                                      Conservative
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Pred. No.
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ADB54361 standard; DNA; 19

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short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anorectic; antidiabetic; antiinflammatory; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel method for detecting and differentiating between colon cell proliferative disorders associated with at least one gene or its regulatory regions. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least one reagent or a series of reagents, where the reagent or series of reagents, distinguishes between methylated and non methylated CpG dinucleotides within the target nucleic acid. The molecules of the invention demonstrate cytostatic activity whilst the method may useful for detecting and differentiating between colon cell proliferative disorders, including cancers such as colon adenoma and colon carcinoma. The PNA (peptide nucleic acid)-oligomers are useful as probes for determining cytosine methylation state or single nucleotide polymorphisms. The current sequence is that of the PCR primer of the properties with the courtes to the polymorphisms.
                                                            Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:171.
                                                                                                29-JAN-2004
                                                                                                                              ADE29549,
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Rujan T, Schmitt A;
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invention which
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associated with
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                                                                                                                                                                                                                                                                                                                      Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                 amplify the genomic DNA region.
                                                                                                                                                                                                                                                                                                                                                                  G; 5 T; 0
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                                                                                                                                                                                                                                                                                                                                                                    U; 0 Other;
                                                                                                                                                                                                                                                                                                                      1.2e+05;
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                                                                                                                                                                                                                                                                                                                                   Length 19;
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SAKAKU
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ADE29386
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                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiasthmatic, immunosuppressive, antibacterial, antirheumatic, antipacterial, antirheumatic, antipacriatic and gastrointestinal activities. The MAPK sinks can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes types I and II; a wide range of tumours, and inflammatory diseases (asthma, septic shock, rheumatoid arthritis, psoriasis and inflammatory bowel disease). They can also be used for drug screening; diagnosis; target identification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide studying screening). The present sequence represents a MAPK sink which is used the property of the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is used to the sequence represents a map which is the sequence 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 genetic engineering; pharmacogenomic; gene mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; SEQ ID NO 171; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein kinase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-FEB-2002;
11-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of a mitogen-activated protein kinase (MAPK) genes by RNA interference. Also described: (1) a method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-JAN-2003;
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                                                                                                                   ADE29386 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2003-689980/65.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                        Conservative
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2003US-0440129P.
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                                                                                                                   RNA;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                      Length 19;
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29-JAN-2004

(first

RESULT 23 AAQ52874 ID AAQ9

AAQ52874

standard; RNA;

20 ₽P 片 S

Matches Query Match Best Local :

11;

Conservative

Similarity

2.0%; 91.7%;

Score 12; Pred. No. Mismatches

DB 9; .2e+05

Length 19

0

Gaps

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зві есеестесьссе зэг

GCGGCUGCACCG

12

Sequence 19

BP; 1 A; 7 C; 10 G; 0 T; 1 U; 0 Other;

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29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                              have cytostatic, anorectic, antidiabetic, antiinflammatory, antiasthmatic, immunosuppressive, antibacterial, antirheumatic, antiarthritic, antipsoriatic and gastrointestinal activities. The MAPK siNAs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obesity diabetes types and II; a wide range of tumours, and inflammatory diseases (asthma, septic shock, rheumatoid arthritis, psoriasis and inflammatory bowel disease). They can also be used for drug screening; diagnosis; target identification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK siNA which is use in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mitogen-activated protein kinase; MAP kinase; MAPK; RAPK; RAM interference; mitogen-activated protein kinase; MAP kinase; MAPK; RAPK; RAM interference; cytostatic; anorectic; antidiabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antirheumatic; antiarchritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour;
                                                                                                                                                                                                                                                                   modulating expression of MAPK genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) vectors that express siNA and cells containing these vectors. MAPK siNAs
                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; SEQ ID NO 8; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid, useful e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcswiggen
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06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a short interfering nucleic acid (siNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis of cancer,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genetic engineering; pharmacogenomic; gene mapping;
                                                                                                                                                                                                                                                                                                                                                    that downregulates expression of a mitogen-activated protein kinase (MAPK) genes by RNA interference. Also described: (1) a method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SIRN-) SIRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            downregulates expression of mitogen-activated
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                                                      which is used
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sequences for enzymatic RNA molecules. The NA molecules are gene target. They also have enzymatic activity, in that they specifically cleave RNA in the target. The ERMs interfere with viral replication and therefore have anti-viral properties. They can be used to attenuate viruses to be used in vaccines. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)
     on 25-MAR-2003
PI field.)
                                                                                                                                                                                   Enzymatic RNA molecules
                                                                                                                                                                                                            WPI; 1993-386599/48
                                                                                                              The sequences
                                                                                                                                         Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA; enzyme; enzymatic RNA molecule; ERM; cleave; RNA; mRNA; HnRNA; picornavirus; HIV; immunodeficiency virus; hepatitis B virus; HBV; papilloma virus; HPV; Epstein-Barr virus; EBV; TCLV; T-cell leukaemia virus; hepatitis C virus; HCV; cytomegalovirus; influenza virus; HSV; herpes simplex virus; vector; immune response;
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26-MAY-1994
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                                                                                                                                                                                                                                                                               RIBOZYME PHARM
                                                                                                                                        Fig 13; 287pp; English.
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                                                                                                         (AAQ52824-Q52890) are pref. Cytomegalovirus target
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(first entry)
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92US-00987130.
92US-00987133.
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92US-00884073.
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92US-00882922.
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                                                                                                                                                                              inhibit viral replication,
                                                                                                                                                                                                                                                    Macejak DG,
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RESULT 238
AAQ94258/c
ID AAQ942
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AC AAQ942
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AC AAQ942
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DT 05-DEC
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DT 05-DEC
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primer; PCR; polymerase chain reaction; SV40; large T antigen;
mouse hepatoma cell; animal model; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                             The primers (AAQ55811-841) are used to detect various types of hepatitis C virus. The primers are made from oligo DNA fragments selected from specific hepatitis C virus subtypes. The primers can be used to in a one step PCR reaction which can determine the subtypes of a large number of
                                        Antisense primer to amplify 429 bp SV40 large T antigen gene fragment
                                                                           05-DEC-1995
                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detection of type C hepatitis virus - using reaction with mixed primer set.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Page 2; 7pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-037380/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HCV; hepatitis C virus; detection; primer; PCR; mixer polymerase chain reaction; DNA polymerase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HCV detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ55841 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SAYA/) SAYAMA
                                                                                                                                                                                                                                              36 TTACCAATTCAA 47
                                                                                                                                                                                                                                                                          Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 UGCUACCGCGUC
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                                                                                                                                      standard;
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                                                                                                                                                                                                                                                                                                                                   BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;
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ilarity 75.0%;
Conservative
                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                      DNA;
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C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Japanese
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12; DB 2;
Pred. No. 1.2e+05
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                                                                                                                                                                                                                                                                                       DB 2; Lo., 1.2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      one step DNA polymerase chain
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Matches
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                                                                                                   24-MAR-1994;
05-AUG-1994;
04-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ94257, to amplify a region from nucleotides 1571 to 2009 of the SV40 large T antigen gene. A fragment of 429 by was generated. The fragment is used in the invention, a mouse hepatoma cell line which may metastasise specifically in the liver and contains an albumin promoter gene and a SV40-T gene. The cell line is useful for the production of animal models for the study of cancer and screening of anti-cancer agents
                                                             Burmer
                                                                                                                                                                                                                                    Human herpes virus-6; HHV-6; multiple sclerosis; genetic marker; MDBP; internal primer MSHind9; diagnosis; ss.
                                                                                                                                                                                                                                                                                                            AAQ91041;
                                                                                                                                                                                                                                                                                                                              AAQ91041 standard; DNA;
                                                                               (PATH-) PATHOGENESIS CORP
                                                                                                                                 05-NOV-1993;
                                                                                                                                                      04-NOV-1994;
                                                                                                                                                                          11-MAY-1995
                                                                                                                                                                                              WO9512313-A1.
                                                                                                                                                                                                                                                                HHV-6 associated MS genetic marker MDBP internal primer MSHind9.
                                                                                                                                                                                                                                                                                        30-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 5 A; 2 C; 4 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AD094257, to amplify a region from nuclectides 1571 to 2009 of the SV40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse hepatoma cell line - useful in animal model systems
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                                                            Challoner PB,
                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                      (first
                                                                                                 93US-00149176.
94US-00218029.
94US-00287942.
94US-00334482.
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                                                                                                                                                     94WO-US012655
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thes 0;
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Treatment of human herpes-virus-6-associated multiple sclerosis - u an antiviral agent, e.g. a nucleoside analogue, administered to the

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                                                                     CC Rapid and cost effective diagnosis of disease-associated mutations in the CC p53 gene is achieved by employing a selected number of diagnostic tools, CC in a hierarchy of increasing accuracy and cost per tool, in which each CC tool detects essentially no false positives. Tests that may be employed, CC in order of increasing accuracy and cost are: (a) immunoassays; (b) DNA CR fragment length/quantity analysis; and (c) DNA sequencing of regions CC most likely to harbour point mutations. AATO8645-66 are primers used in CC DNA fragment length/quantity analysis. The amplification of the eleven CC exons is advantageously carried out in 3 multiplex pools, the members of CC a pool selected because they all use the same hybridisation temperature CC and none of the expected fragment lengths will overlap in an CC electrophoresis gel. One of each pair of primers is labeled at the 5' end with an identifiable marker such as fluorescein, rhodamine or cyanine. The present sequence is used with AATO8654 to amplify a 247 bp fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Testing for disease-associated p53 gene mutation(s) using a hierarchy assay techniques - e.g. immunoassay, DNA amplification and DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-JUL-1994;
14-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 18; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JUL-1995;
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quantitative assay; fragment length; DNA sequencing; p53; mutat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (VISI-) VISIBLE GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JAN-1996
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95US-00388381.
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Query Match Best Local :

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                                                      Query Match
Best Local S
Matches 12
                                                                                                                                               Rapid and cost effective diagnosis of disease-associated mutations in the possible problem of participation of problem of participations in the present sequence is achieved by employing a selected number of diagnostic tools, in a hierarchy of increasing accuracy and cost per tool, in which each tool detects essentially no false positives. Tests that may be employed, in order of increasing accuracy and cost are: (a) immunoassays, (b) DNA fragment length/quantity analysis, and (c) DNA sequencing of regions most likely to harbour point mutations. Alamonation of the erven exons is advantageously carried out in 3 multiplex pools, the members of a pool selected because they all use the same hybridisation temperature and none of the expected fragment lengths will overlap in an electrophoresis gel. One of each pair of primers is labeled at the 5' end with an identifiable marker such as fluorescein, rhodamine or cyanine. The present sequence is used with AATO8651 to amplify a 268 bp fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                             Testing for disease-associated p53 gene mutation(s) using a hierarchy of assay techniques - e.g. immunoassay, DNA amplification and DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-097638/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diamandis E,
                                                                                                                                                                                                                                                                                                                                                                          Claim 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (VISI-) VISIBLE GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JUL-1994;
14-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer; PCR;
quantitative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT08652;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9601909-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer P53-3X5MP for p53 gene exon 5 amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT08652 standard; DNA; 20 BP.
                         510 regrererecae 521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  510 TCGTCTCTCCAG 521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18 TCGTCTCTCCAG 7
ø
                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                        Page
                                                                                                           BP; 2 A; 9 C; 4 G; 5 T; 0 U; 0 Other;
                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymerase chain reaction; hierarchy; immunoassay; assay; fragment length; DNA sequencing; p53; mutation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dunn JM,
                                                                                                                                                                                                                                                                                                                                                                        21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94US-00271946.
95US-00388381.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95WO-US008605
                                                                                                                                                                                                                                                                                                                                                                     44pp; English
                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.0%;
20
                                                                  2.0%; Score 12; DB 2; Length 20; 100.0%; Pred. No. 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stevens
                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12;
Pred. No.
                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2; hc. o. 1.2e+05; 0;
                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                     0;
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                                                  Gaps
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                                                   0,
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RESULT

RESULT 243
AAT99863
ID AAT998
XX
AC AAT998
XX
DT 07-MAY

AAT99863 standard; DNA;

20 ₽P.

07-MAY-1998

(first

entry)

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AAT99864/c

ID AAT99864;

XX

AC AAT99864;

XX

DT 07-MAY-1998 (first entry)

XX

PCR primer for exon 6 of p53 gene.

XX

PCR primer; amplify; pathogen id

KW nucleic acid analysis; microorga

KW HLA type determination; p53 gene

XX

PN W09741259-A1.

XX

VISIBLE GENETICS INC.

PR 19-JUL-1996; 96US-00684498.

PR 27-FEB-1997; 97US-00807138.

XX

VISIBLE GENETICS INC.

XX

VISII-) VISIBLE GENETICS INC.

XX

Nucleic acid sequence determinate extension products, which are in prime for mucleotide in nucleotide sequence represents a prime extension products, which are in prime for mucleotide in the region of inte sequence can be used in the meth companies combining the sample with the companies of inte produced, characterised in that region of inte produced, characterised in that companies combining the sample with the companies of inte produced, characterised in that companies of inte produced in that companies of interproduced in that produced in that produced in the produced in that p
                                                                                                                                                                                                                                                                                                                                               This sequence represents a primer for exon 6 of the p53 gene. This sequence can be used in the method of the invention for determining the position of at least one selected species of nucleotide, in a region of interest, in a target nucleic acid polymer, in a sample. The method comprises combining the sample with a reaction mixture to synthesise chain extension products indicative of the positions of the species of nucleotide in the region of interest and evaluating the products produced, characterised in that the sample, which is combined with the reaction mixture, and contains target and non-target nucleic acid reaction mixture, and contains target and non-target nucleic acid colymers in natural abundance. The method can be used to detect mutations, particularly mutations of medical significance, in samples derived from a human patient, animal, plant or microorganism, determine HLA type ancillary to transplant procedures, detect and identify microorganisms, particularly pathogenic microorganisms, in a sample and in in situ sequencing reactions to produce sequencing fragments in a contains of histological specimen
                                                                                                                                           Matches
                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 4 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid sequence determination - comprising synthesising chain extension products, which are indicative of positions of selected species of nucleotide in nucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 4; Page 19; 69pp; English.
                                                                                                                                                                             Local
                                                               510 TCGTCTCTCCAG 521
18
                                                                                                                                   l Similarity
12; Conserv
                                                                                                                                       Conservative
                                                                                                                                                                             100.0%;
                                                                                                                                                                                                               2.0%;
                                                                                                                                       0; Mismatches
                                                                                                                                                                             Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Larson MT,
                                                                                                                                                                         1.2e+05
                                                                                                                                                                                                           DB 2;
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                                                                                                                                                                                                               Length 20
                                                                                                                                       Indels
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AAT79479/
IID AAT7
XX
AC AAT7
XX
AC AAT7
DT 22-O
XX
DE DNA
XX
XX
Aden
KW 11ga
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                                                                                                                                                                                          RESULT 244
                                                                                                                                                                                                                                                                                                                      Best
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                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
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19-JUL-1996;
27-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          interest, in a target nucleic acid polymer, in a sample. The method comprises combining the sample with a reaction mixture to synthesise chain extension products indicative of the positions of the species of nucleotide in the region of interest and evaluating the products produced, characterised in that the sample, which is combined with the reaction mixture, and contains target and non-target nucleic acid polymers in natural abundance. The method can be used to detect mutations, particularly mutations of medical significance, in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid sequence determination - comprising synthesising chain extension products, which are indicative of positions of selected species of nucleotide in nucleotide sequence.
Adenosine; adenosine-5'-phosphate; adenosine triphosphate; ATP; binding; ligand; purification; reagent; isolation; determination; subcellular localisation; catalyst; assay; SELEX;
                                                                                                                                                                                                                                                                                                                                                                                                                              derived from a human patient, animal, plant or microorganism, HLA type ancillary to transplant procedures, detect and ident: microorganisms, particularly pathogenic microorganisms, in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence represents a primer for exon 5 of the p53 gene. This sequence can be used in the method of the invention for determining the position of at least one selected species of nucleotide, in a region of in a target nucleic acid polymer, in a sample. The method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer; amplify; pathogen identification; mutation detection; nucleic acid analysis; microorganism characterisation; human;
                                                                                                                                                            AAT79479 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 19; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9741259-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                               22-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                   histological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Leushner J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HLA type determination; p53 gene exon 5; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer
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                                                                                                                                                                                                                                                                                                                                                                                                                 in situ sequencing reactions to produce sequencing fragments in a
                                                                                                                                                                                                                                                                        510
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                                                                                                                                                                                                                                                                                                     l Similarity
12; Conserv
                                                                                                                                                                                                                                        regrereredad 20
                                                                                                                                                                                                                                                                      regrererecae 521
                                                                                                                                                                                                                                                                                                                                                                 BP; 2 A; 9 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hui M,
                                                                                              (first
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96US-00684498.
97US-00807138.
                                                              adenosine or adenosine 5'-phosphate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97WO-US007135
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Б
                                                                                              entry)
                                                                                                                                                                                                                                                                                                                 2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              p53
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                                                                                                                                                                                                                                                                                                     ٥,
                                                                                                                                                                                                                                                                                                                   Score 12; DB 2; 1
Pred. No. 1.2e+05
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                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lacroix J,
                                                                                                                                                                                                                                                                                                                                   Length 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shipman
                                                                                                                                                                                                                                                                                                                                                                                                                                 in a sample and
                                                                                                                                                                                                                                                                                                                                                                                                                                              inism, determine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT91100 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 7 A; 1 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        stability and synthesis cost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      phosphate - useful as purilication reagent, or rot vectors adenosine tri:phosphate subcellular localisation in vivo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single stranded DNA molecule, which binds adenosine or adenosine-5'-phosphate - useful as purification reagent, or for determination of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-288574/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Szostak JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Systematic Evolution of Ligands by Exponential enrichment;
                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; Col 63-64; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JAN-1995;
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                                                                                                                                                                                                                                                           LAMAN; lysosomal alpha-mannosidase; alpha-mannosidosis; diagnosis; screening; genetic test; PCR; primer; RFLP;
                                                                                                                                                                                                                                restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is an adenosine or adenosine-5'-phosphate (ASP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16 CTTCCTCCGCTA 5
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                                                                                                                                                                                                                                                                                                                                                       lysosomal alpha-mannosidase (LAMAN) gene PCR primer mp5UT1F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                            screening; generations the screening servers to serve servers to server to servers to server to servers to server to serv
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                                                                                                                                                                                                                                                                                                                                                                                                                           (first
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                                                                                                                                                                                                                             polymorphism;
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Pred. No.
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. 1.2e+05
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24-JUL-1997

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RESULT 246
AAT99833
ID AAT998
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Best Local
                                                                                               01-MAY-1996;
19-JUL-1996;
                                                                                                                               29-APR-1997;
                                                                                                                                                     06-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Forward primer mp5UTIF is based on a 1500 bp amplicon produced from bovine fibroblast genomic DNA using primers (see AAT91098-99) based on the bovine alpha-mannosidase (LAMAN) gene (see AAT91086). It was used with reverse primer mp262 (see AAT91101) to obtain an 800 bp RT-PCR product that constituted a 5' part of LAMAN cDNA. This was combined with a previously obtained PCR cDNA fragment (see AAT91096-97) to produce a full-length clone (see AAT91086) for LAMAN (see AAW26682). Mutations in
                                                   Leushner J,
                                                                                                                                                                           WO9741258-A1
                                                                                                                                                                                                                                         PCR primer; amplify; p53 gene exon 5; multiplex amplification nucleic acid analysis; microorganism characterisation; human;
                                                                                                                                                                                                                                                                                                   07-MAY-1998
                                                                                                                                                                                                                                                                                                                                             AAT99833 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Diagnosis and screening for bovine alpha-mannosidosis - by detecting mutation(s) in alpha-mannosidase gene, also nucleic acid encoding the enzyme and derived oligo:nucleotide primers.
                                                                                                                                                                                                                                 mutation detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the LAMAN gene cause bovine alpha-mannosidosis, and can be detected using claimed PCR primers (see AAT91088-93)
                                                                        (VISI-) VISIBLE GENETICS INC.
                                                                                                                                                                                                 Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the LAMAN
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12; Conser
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DZIEGLEWSKA
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                                                                                                                                                                                                                                                                             exon 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 22; 85pp; English.
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                                                  Hui M,
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                                                                                              96US-00640672.
96US-00684498.
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Pred. No.
                                                   Larson
                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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Analysing nucleic acid amplification reaction

containing sample - comprises performing multiplex and reacting amplified fragments in sequencing

This sequence represents a primer for exon 6 of the p53 gene. This sequence can be used in the method of the invention for analysing a nucleic acid containing sample. The method comprises performing a multiplex amplification reaction on the nucleic acids in the sample using amplification primer pairs, one pair for each region to be analysed, to produce a mixture of amplified fragments, and determining the sequence of

Analysing nucleic acid containing sample - comprises performing multiplex amplification reaction and reacting amplified fragments in sequencing

Example

4

Page 18;

37pp; English

reaction mixture

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RESULT 247
AAT99834/c
ID AAT998
XX AAT998
XX AAT998
XX Primer
XX PCR pr
KW nuclei
KW nuclei
KW nuclei
KW nuclei
XX Synthe
OS Homo g
XX HOMO 14
XX 19-JUI
XX 19-JUI
XX 29-APF
XX 01-MAY
PR 19-JUI
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PT ampli;
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19-JUL-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a primer for exon 5 of the p53 gene. This sequence can be used in the method of the invention for analysing a nucleic acid containing sample. The method comprises performing a multiplex amplification reaction on the nucleic acids in the sample using amplification primer pairs, one pair for each region to be analysed, to produce a mixture of amplified fragments, and determining the sequence of at least one species of amplified fragment, characterised in that the sequence is determined by combining the mixture of amplification of fragments with a sequencing reaction mixture for the production of sequencing fragments, and evaluating the sequencing fragments produced. The method can be used to analyse regions in the nucleic acids in the sample for the presence of mutations, or detect and type microorganisms. The method directly performs sequencing reactions on complex DNA mixtures
                                                                                                                                                                                                                                                                                                                                                                                          WO9741258-A1
                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; amplify; p53 gene exon 6; multiplex amplification nucleic acid analysis; microorganism characterisation; human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mutation detection;
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12; Conserv
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96US-00684498.
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Pred. No.
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1.2e+05;
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RESULT 248
AAV47975
ID AAV479
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AC AAV479
XX
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Best Local
The oligonuclectides which specifically hybridise to B7 modulate its expression (and thus T cell activation and proliferation). This is particularly useful for treatment and prevention of inflammation and autoimmune diseases, e.g. asthma, (juvenile) diabetes, myasthenia gravis, Grave's disease, rheumatoid arthritis, allograft rejection, psorilasis, (gystemic) lupus erythematosus, multiple sclerosis, contact dermatitis, rhinitis, allergy, cancer and metastases. The oligonucleotides may also be used to manipulate T cell activation ex vivo; to determine or detect B7 protein expression; for diagnosis; as assay and purification reagents, and to study physiological roles of B7 proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           at least one species of amplified fragment, characterised in that the sequence is determined by combining the mixture of amplification fragments with a sequencing reaction mixture for the production of sequencing fragments, and evaluating the sequencing fragments produced. The method can be used to analyse regions in the nucleic acids in the sample for the presence of mutations, or detect and type microorganisms. The method directly performs sequencing reactions on complex DNA mixtures
                                                                                                                                                                                                                                                                                                     New oligo:nucleotide(s) that modulate expression of B7 proteins - used for, e.g. controlling activation and proliferation of T cells, particularly for treatment, diagnosis and prevention of inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Vickers TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PHARM INC.
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/note=
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"Phosphorothioate linkages"
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                                                                                                                                                                                                                                                           English
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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hes 0;
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TIGGTIATCITC 148

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12;

Conservative

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Similarity

2.0%;

Score 12; Pred. No. Mismatches

DB 2; Lo., 1.2e+05;

Length 20

Indels

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Gaps

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Query Match Best Local

Sequence

20

BP; 5 A; 4 C; 3 G; 8 T; 0 U; 0 Other;

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RESULT 249
AAV5598
ID AAV559
XX AAV559
XX AAV559
XX AAV559
XX O4-DEC
XX GTPCH-
XX Select
KW THB; G
KW Inflam
KW Inflam
KW Inflam
KW INFO
OS HOMO 6
XX WO9835
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                                                    CC sequences shown in AAV55996 to AAV56003 represent primers used for RT-PCR CC amplification during the course of the invention. The invention provides a method for identification of selective inhibitors of nitric oxide CC synthases (NOS) which comprises determining inhibition of activity of NOS in the presence of a candidate compound and two different concentrations of tetrahydrobiopterin (THB). Inhibitor of GTP cyclohydrolase 1-mediated synthesis of THB can be identified by contacting GTP cyclohydrolase 1 which a candidate compound and determining inhibition of the enzyme CC with a candidate compound and determining inhibition of the enzyme CC identified by the methods may be used in treatment of conditions in which the production of nitric oxide (NO) is implicated, such as septic shock, CC asthma, arthritis, inflammatory bowel disease, heart failure or acute systemic inflammation. The inhibitors may also be useful in treatment of neurological disease states in which neuronal NOS has been implicated, CC e.g. stroke, dementia or Parkinson's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTPCH-1 mRNA amplifying RT-PCR primer igc1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               determining
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identification of selective inhibitors of nitric:oxide synthases determining inhibition of enzyme activity in presence of candidations of tetra:hydro:biopterin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UNIO ) UNIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9835055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          stroke; RT-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rccreekecccc 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCCTGGAGCCCC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bhagat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COLLEGE LONDON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97GB-00002312.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GTPCH-1; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vallance PJT, Hingorani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
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Pred. No. 1.2
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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1.2e+
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 251
AAZ31857
ID AAZ318
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AC AAZ318
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DT 24-JAN
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DE PCR pr
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                                                                                                                                                                                                                                         Best Local Similarity Matches 12; Conserv
                                                                                                                                                                                                                                                         Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a method for measuring the resistance of p53 mutant cancer cells to the cytotoxic effects of chemotherapeutic agents by testing a sample comprising p53 mutant cells or an extract from p53 mutant cells for the abundance of cyclin protein D1. AAX9036 to AAX90373 represent reverse transcription PCR primers used to amplify the human p53 gene. The method can be used to predict the response of human cancer cells to anticancer therapy agents which can be used to select the most appropriate therapy for patients suffering from cancer. High cyclin D1 levels or high cyclin D1 expression together with p53 mutation is strongly associated with resistance to cis-diaminedichloroplatinum (CDDP)
                                                                                                                                                                                                                                                                                                                                   in human cancer cells. The test may be used to detect resistance to other cytotoxic agents such as etoposide and indicate whether radiation may be a viable alternative to CDDP or if other cytotoxic agents would be more suitable, e.g. may suggest that Taxol should be considered as an alternative therapy as it may not be sensitive to a combination of p53 mutation and cyclin D1 protein overexpression
                                                                                                                                                                                                                                                                                                       Sequence
PCR primer for human G-alpha-13 coding sequence
                                  24-JAN-2000
                                                                                               AAZ31857 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 13; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Measuring resistance of p53 mutant cancer cells to cytotoxic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-422070/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Warenius HM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GB2334577-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human p53 gene reverse transcription PCR primer exon 5 antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX90365 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYLI-) UNIV LIVERPOOL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                          510 TCGTCTCTCCAG 521
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                                                                                                                                                                                                                                                                                                       20 BP; 2 A; 9 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                            TCGTCTCTCCAG 20
                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reverse transcription; PCR primer; resistance; mutant; lin D1 protein; chemotherapy; cytotoxic; ss.
                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98GB-00003446
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                                                                                                BP.
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                                                                                                                                                                                                                                                         Score 12;
Pred. No.
                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                      DB 2; Lo
                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                         Indels
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AAV80729/
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                                                                                                                                                                                                                                                                                                                                 RESULT 252
                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                         Porcine; wild boar; meishan; pietrain; large white; hampshire; duroc; differentiation; breed origin; alpha-MSHR; coat colour; stock purity; alpha melanocyte-stimulating hormone receptor; KIT; PCR primer; ss.
                                                                                                                                                                                                                                                                                                       729/c
AAV80729 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence represents a PCR primer for DNA encoding the human 13 protein. The invention relates to an antisense compound of 8 t nucleobases in length, that inhibits the expression of the human 13. The antisense compound is useful for treating an animal, part humans, having or being prone to a disease or condition associate the expression of G-alpha-13, such as cancer
         Andersson L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 14; Col 37; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G-alpha-13; human; inhibitor; cancer; antisense compound; therapy; PCR primer; ss.
                                                     30-MAY-1997;
31-JAN-1998;
                                                                                       27-MAY-1998;
                                                                                                             03-DEC-1998
                                                                                                                                    WO9854360-A1
                                                                                                                                                          Sus scrofa.
                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                       Alpha melanocyte-stimulating hormone receptor PCR primer EPIG13.
                                                                                                                                                                                                                                                               26-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense compound inhibiting expression of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-633376/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                               (PIGI-) PIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5981732-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
les 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                        319 AGGATCTTCACC 330
                                                                                                                                                                                                                                                                                                                                                                   9
                               IMPROVEMENT CO UK LTD
                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHARM INC
                                                                                                                                                                                                                                                             (first entry)
         Kijas J,
                                                     97GB-00011214.
98GB-00001990.
                                                                                       98WO-GB001531.
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                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                     2.0%;
        Giuffra
                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to a disease or condition associated with such as cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                          No.
         Evans
                                                                                                                                                                                                                                                                                                                                                                                                                        1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2;
         원
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 20
         Wales
                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G-alpha-13
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          Plastow
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G-alpha-
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Differentiating products from different of alleles of breed-determinant genes, a

t animal breeds - by at the nucleic acid

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protein

the analysis

WPI; 1999-070222/06

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   products according to breed origin; (b) determining or testing the breed corigin of a product; or (c) validating an animal product. The method comprises analysing a sample of the product for the allele(s) of at least come breed-determinant (BD) gene. The present invention also describes: (1) methods for determining the coat colour genotype of a pig by come receptor (alpha-MSHR) gene; (ii) the anino acid sequence of an colour manner receptor (alpha-MSHR) gene; (ii) the anino acid sequence of an colour for the protein; (iii) detecting which microsatellites (or other linked comarker alleles), linked to the alpha-MSHR gene, or particular alleles of it, are present; and (iv) analysing nucleic acid to determine if the KIT gene carries a polymorphism associated with the Belt genotype. The main commanls, especially pigs. Particular applications are confirming stated corigin of meets; in quality control; for maintaining stock purity, and in coreding programmes (to confirm particular crosses). The method ce inexpensively. The process can be made quantitative. The present sequence conly very small samples and many samples can be screened quickly and convery control controls for maintaining stock purity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 12
               Diagnosis of cancerous or pre-cancerous cyclin-dependent kinases 1 and 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
                                                         WPI; 1999-432548/37
                                                                                                                                                                        18-FEB-1998;
                                                                                                                                                                                                      25-AUG-1999
                                                                                                                                                                                                                                  GB2334578-A
                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                          cyclin-
                                                                                                                                                                                                                                                                                                                      Human;
                                                                                                                                                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                                                                                               24-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                            AAX90393
                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX90393 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A method has been developed for: (a)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 22; Page 70; 101pp; English.
                                                                                                                  (UXLI-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           243
                                                                                                                                                                                                                                                                                                                                                   p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                         p53; reverse transcription; PCR primer; cancer; diagnosis; mutant;
dependent kinase; CDK; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      . Similarity 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CACCICCIGGAG 254
                                                                                                                                                                                                                                                                                                                                                   gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                  LIVERPOOL.
                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                     Seabra L;
                                                                                                                                                                                                                                                                                                                                                   reverse transcription PCR primer exon 5 antisense
                                                                                                                                           98GB-00003447
                                                                                                                                                                         98GB-00003447
                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA; 20
                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                        ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       differentiating animals and animal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.2e+05;
hes 0;
                           cells by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 20;
                           monitoring
                            the levels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                  18-FEB-1998;
18-FEB-1998;
05-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a method for the diagnosis of a cancerous or pre-cancerous state from the co-elevation of cyclin-dependent kinase 1 (CDK1) and CDK4 levels. The method may be used for the clinical diagnosis of cancerous or pre-cancerous cells. In addition the combination of targets may be used to screen for drugs that may specifically act on cancer cells. The combination of CDK1, CDK4 elevation and p53 mutation in combination form a complex target that is likely to be specific for cancerous cells. AAX90388 to AAX90401 represent reverse transcription PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human;
signal
                                        The present invention describes a method for the determination of sensitivity of cancer cells to anti-cancer agents by measuring the mutational status, expression and /or function of signal transduction factors. The method, by measuring the resistance of cells to anti-cancer agents, is useful for selecting the most appropriate therapy for patients suffering from cancer. AXY0374 to AXY0377 represent reverse transcription PCR primer for the human p543 gene, used in an example from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A;
                                                                                                                                                                                    Determination of sensitivity of cancer cells to anti-cancer agents
                                                                                                                                                                                                                 WPI; 1999-422071/36
                                                                                                                                                                                                                                                                                                                                                                           03-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                     GB2334579-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human p53 gene reverse transcription PCR primer exon 5 antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX90379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX90379
                                                                                                                                                       Example 1; Page 18; 46pp; English.
                                                                                                                                                                                                                                              Warenius
                                                                                                                                                                                                                                                                                                                                                                                                       25-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer for the human p543 gene, used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bxample;
                            the present invention
                                                                                                                                                                                                                                                                         (UYLI-) UNIV LIVERPOOL. (THER-) THERYTE LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        510 regrererecae 521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        p53; reverse
transduction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              φ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
12; Conserv
                                                                                                                                                                                                                                              E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOGTOTOTOCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 12; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
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98GB-00003447.
98GB-00012151.
                                                                                                                                                                                                                                                                                                                                                                           98GB-00014545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transcription; PCR primer; cancer;
factor; mutant; cell cycle; apoptos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 C; 4 G; 5 T; 0 U; 0 Other;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chemotherapy,
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Sequence 20

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RESULT 2

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RESULT 256
AAX79784
ID AAX79:
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AC AAX79:
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DT 17-AUC
                                                                                                                                                                                                            Query Match
Best Local S
Matches 12
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                 This invention describes novel antisense oligonucleotides against thioredoxin and thioredoxin reductase gene which have cytostatic activity and are useful for inhibiting tumour growth and metastasis in mammals. They may also be used as hybridization probes to detect the presence of the thioredoxin and thioredoxin reductase mRNAs in mammalian cells. They may also be used as molecular weight markers. The antisense oligonucleotides are nuclease resistant due to the presence of phosphorothioate internucleotide linkages. AAZ00478-Z00503 represent oligonucleotide primers capable of binding to human thioredoxin mRNA
      17-AUG-1999
                                     AAX79784;
                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense oligonucleotides against thioredoxin and thioredoxin reductase genes, useful for inhibiting tumor growth and metastasis.
                                                                   AAX79784 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 18; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-469328/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thioredoxin; thioredoxin reductase; human; antisense; primer; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ00491 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                     442 TGGAATACTTTT 453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                510 regrererecae 521
                                                                                                                                                                                                            12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENESENSE TECHNOLOGIES INC.
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                                                                                                                                                                                                                                                                         BP; 5 A; 4 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Young AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumour growth inhibitor; detection; nuclease resistant,
      (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-0073196P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linkage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA binding antisense oligonucleotide 2614.
                                                                   DNA;
    entry)
                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.0%;
                                                                                                                                                                                                                                           2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lee YS;
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                                                                   ВP
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                                                                                                                                                                                                                         Score 12; DB 2; L
Pred. No. 1.2e+05;
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Pred. No.
                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                          Length 20;
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                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                         Gaps
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RESULT 257
AXX92037
ID AAX920
XX AAX920
XX 13-SEP
XX PCR pr
XX Respir
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XX Synthe
OS Chlamy
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PR 21-NON
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   21-NOV-1997;
04-NOV-1998;
                                               20-NOV-1998;
                                                                                                                                                              Synthetic
                                                                                                                                                                                                         Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 10 A; 4 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence represents a PCR primer that can be used in the method of the invention. The method is for genetic diagnosis using human mitochondrial DNA where there is at least one base replacement from among the following five replacements: the 3010th base is changed from guanine to adenine; the 4883rd base from cytosine to thymine; the 5178th base from cytosine to adenine; the 8414th base from cytosine to thymine; and the 1468th base from cytosine to thymine. The method can be used for diagnosing the probability of contracting adult diseases. A confirmation of base replacement can give a diagnosis of the level of probability of contraction of adult diseases
                                                                                03-JUN-1999
                                                                                                               WO9927105-A2
                                                                                                                                                                                          neutralising epitope;
                                                                                                                                                                                                                                                         PCR primer used to amplify an ORF of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                         13-SEP-1999
                                                                                                                                                                                                                                                                                                                       AAX92037;
                                                                                                                                                                                                                                                                                                                                                      AAX92037 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 6; 15pp;
                                                                                                                                            Chlamydophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      base replacements
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genetic diagnosis using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer; human; mitochondrial DNA; genetic diagnosis, adult disease contraction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer H16016 for mitochondrial DNA analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12;
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                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                            pneumoniae.
                                                 98WO-IB001890.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                     citis media; erythema nodosum; pharyngitis; vaccine;
PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.0%;
                                                                                                                                                                                                                                                                                                                                                      20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Japanese.
                                                                                                                                                                                                                                                                                                                                                      BP.
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Pred. No.
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RESULT 258
AAC61817/c
ID AAC618
XX AAC618
XX AC618
XX AC6
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Matches 12; Conserv
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; Fas; Apo-1; antisense compound; Fas ligand; Fap-1; hepatitis; Fas associated protein 1; protein tyrosine phosphatase; cancer; autoimmune disease; inflammatory disease; lymphoma; phosphorothioate;
     Dean NM,
                                                                                                                                                                                                                         19-OCT-2000
                                                                                                                                                                                                                                                                               WO200061150-A1
                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               misc_feature
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                                                                                                                                                                   10-APR-2000; 2000WO-US009540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
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                                                           ISIS
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     Marcusson EG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; 1912pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C. pneumoniae
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                                                              PHARM
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                                                                                                              99US-00290640
                                                                                                                                                                                                                                                                                                                                                                                    /note= "2'-methoxyethoxy residues"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "contains phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                           *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                          *tag=
                                                              INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.0%;
                                                                                                                                                                                                                                                                                                                               "2'-methoxyethoxy residues'
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; Pred. No. 1.2;
0; Mismatches
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1.2e+05;
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RESULT 259
AAA37589
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      untranslated region of nucleic acid molecules encoding Fas, Fas ligand (Fash), or Fap-1 (Fas associated protein 1, protein tyrosine phosphatase). The antisense compounds are used to inhibit the expression of Fas, Fash or Fap-1 in cells or tissues. They are used to treat autoimmune or inflammatory diseases such as hepatitis. They can also be
                                                                                                                                                                                                                                                                                                                                                             paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                            Peptide nucleic acid; PNA; telomerase; ribonucleoprotein inhibitor; neoplasia; neurodegenerative disease; aging; h AIDS; HIV; fungal infection; forensic identification; det
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC61799-C61819 represent antisense oligonucleotides which are directed against nucleic acids encoding human Fas (Apo-1). The specification describes antisense compounds which are targeted to the 5'-untranslated region, translational start site, translational termination region or 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotides for treating hepatitis and colon, livlung cancer are inhibitors of Fas, Fas ligand or Fas associated (Fap-1) expression.
                                                                                       WPI; 2000-292432/25
                                                                                                                                                                 09-APR-1996;
                                                                                                                                                                                            09-APR-1997;
                                                                                                                                                                                                                      04-APR-2000
                                                                                                                                                                                                                                                US6046307-A
                                                                                                                                                                                                                                                                                               misc_feature
                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense sequence #47 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA37589 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 7 A; 4 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         used to treat cancer, especially colon, liver or lung cancer or lymphoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 45; 116pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-628395/60
                                                                                                                                         (TEXA ) UNIV TEXAS SYSTEM
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12; Conserv
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                                                                                                              Piatyszek
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                   96US-00630019
                                                                                                                                                                                           97US-00838545
                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                      /note= "Phosphorothioate internucleotide linkages"
                                                                                                                                                                                                                                                                                       /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA:
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Pred. No.
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                                                                                                               Norton
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1.2e+05;
                                                                                                               JC,
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                                                                                                               Corey
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                                                                                                                                                                                                                                                                                                                                                                              detect; tumour;
                                                                                                                                                                                                                                                                                                                                                                                         hyperplasia;
                                                                                                                                                                                                                                                                                                                                                                                                       enzyme;
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                                                                                                                                                                                                                                                                                                                                                                                                        cancer;
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New peptide nucleic acid (PNA) in mammalian cells is useful as mammalian telomerase.

(A) compounds that inhibit telomerase activity as probes to detect the RNA component of a

Example 1;

C01

29; 45pp;

The present sequence represents an antisense oligonucleotide used as control sequence alongside a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibit elomerase activity. Telomerase is a ribonucleoprotein enzyme

inhibits

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CC affords greater resistance of degradation by proteases or nucleases. The CC therapeutic PNAs may be used for treating disease conditions such as CC cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human CC immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency conditions) and associated pathologies, fungal infections, and other CC syndrome) and associated pathologies, fungal infections, and other CC cytostatic agents, antifungal agents, and other cytotoxic or CC cytostatic agents, antifungal agents, and other cytotoxic or CC cytostatic agents, antifungal agents, and other cytotoxic or CC cytostatic agents. PNA probes are also used for forensic identification CC probes to detect or quantitate polynucleotides having a human telomerase CC RNA (hTR) sequence. PNA probes are also used for forensic identification CC fragment length polymorphism (RFLP) pattern. PNAs are also useful as CC probes to detect the RNA component of amammalian telomerase and as CC probes to detect the RNA component of amammalian telomerase and as CC probes to detect the RNA component of amammalian telomerase and conditions to be detected with increased confidence and CC possibly at an earlier stage, before cells are detected as cancerous CC based on pathological characteristics. The diagnostic and prognostic CC methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the CC cell expresses telomerase activity and its RNA component
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Best Local
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   09-APR-1997;
                                                                       04-APR-2000.
                                                                                                                                                                                                                                                                                                                                                               misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA37584 standard; DNA; 20
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                                                                                                                                               US6046307-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PNA sequence #42 used to inhibit telomerase activity
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12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
97US-00838545
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                                                                                                                                                                                                    /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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Pred. No.
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RESULT 261
AAZ77376/c
ID AAZ773
XX AAZ773
XX AAZ773
XX IO-SEP
XX Human
XX Human
XX Human
XW Human
XW Genomi
XW Genomi
XW Aaplote
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC affords greater resistance of degradation by proteases or nucleases. The CC therapeutic PNAs may be used for treating disease conditions such as CC cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human CC immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency conditions) and associated pathologies, fungal infections, and other CC diseases characterized by abhormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cytotoxic or CC used for molecular diagnostics, labelled PNAs are used as hybridization CC used for molecular diagnostics, labelled PNAs are used as hybridization CC used for molecular diagnostics, labelled PNAs are used as hybridization CC probes to detect or quantitate polymucleotides having a human telomerase CC RNA (hTR) sequence. PNA probes are also used for forensic identification CC fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as CC probes to detect the RNA component of a mammalian telomerase and as CC inhibitors of telomerase activity. The method of the present invention CC allows cancerous conditions to be detected with increased confidence and CC possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic CC methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the CC cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 12
                                                                         Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                       10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                        AAZ77376 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      synthesizes one strand of the telomeric DNA, using as a template an 11 mucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and
                                                                                                                                                                                                Human biallelic marker downstream amplification
                                                                                                                                                                                                                                                                                                         AAZ77376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibit telomerase activity. Telomerase is a ribonucleoprotein enzyme that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Col
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-292432/25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    319 АССАТСТТСАСС 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 5 A; 6 C; 3
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                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12;
Pred. No.
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                                                                                                                                                                                              primer
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                                                                                                                                                                                                   SEQ ID NO:11732
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RESULT 262
AAZ47037
ID AAZ470
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XX AAZ470
XX AATivi
AC Primer
XX Antivi
KW Antivi
KW hepati
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the pharmaceutical states of the characterisation of the pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 6 A; 2 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                     Antiviral; anticancer; antiproliferative; human; interferon-alpha5; hepatic disease; hepatitis C; viral cirrhosis; hepatocellular carcinoma;
                                                                                                                                                                                                                                                                                                                                                      AAZ47037 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the in-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       map
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23-NOV-1998;
                                               13-MAY-1999;
                                                                                                              WO9958143-A1
                                                                                                                                            Homo sapiens
                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                        15-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ65654 to AAZ69578 represent human biallelic markers from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
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                                                                                18-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                    14 CGTCTCTCCAGA 3
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                                                                                                                                                                                          gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                CGTCTCTCCAGA 522
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                                                                                                                                                                                                                                                        for
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                                                                                                                                                                                          expression;
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                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                        human
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98US-0109732P
                  98ES-00001003
                                                 99WO-ES000134
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                                                                                                                                                                                          primer;
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Pred. No.
                                                                                                                                                                                           PCR;
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                                                                                                                                                                                          amplification;
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1.2e+05;
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RESULT 263
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                          chromosome
phenotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method of using interferon-alpha5 or its coding sequence to prepare compositions for treatment of hepatic diseases, e.g. (i) chronic hepatitis C; (ii) cirrhosis of viral origin and (iii) hepatocellular carcinoma. The method restores the level of interferon-alpha5, which is reduced in diseased liver cells, to normal levels. The primers AAZ47036-Z47037 were used initially to detect the level of expression of beta-actin gene in liver tissue by PCR as a control for detecting the level of interferon-alpha or beta. The prime amplify a 314 bp fragment of the human beta-actin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA66151 standard;
                                                                                                                New radiation hybrid map of the dog, Canine familiaris, for e.g. identifying genes implicated in phenotypic and or in genetic diseases and for studying dog pedigrees.
                                                                                                                                                                                                                                                                                                                                                                                Canis familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                       Dog; genome; genomic marker; radiation hybrid map; identification; chromosome location; gene marker; polymorphic microsatellite marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA66151;
The present invention describes a radiation hybrid map of the dog (Canifamiliaris) genome comprising the genome location of a marker selected from AAA6613 to AAA66942. The radiation hybrid map is useful for identifying and localising dog genes, since it covers approximately 80 of the dog genome and provides a dense map integrating different types
                                                                                                                                                                                                                                                                                           15-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating liver diseases with interferon-alphas it, particularly chronic hepatitis C.
                                                                                        Claim 1;
                                                                                                                                                                            WPI; 2000-387821/33.
                                                                                                                                                                                                                                                              13-NOV-1998;
                                                                                                                                                                                                                                                                                                                       25-MAY-2000
                                                                                                                                                                                                                                                                                                                                                    WO200029615-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 11; 36pp; Spanish.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prieto Valtuena J,
                                                                                                                                                                                                                                   (CNRS ) CNRS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genomic marker oligonucleotide sequence
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                                                                                      Page 53; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGAGGATCTTCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGAGGATCTTCA 328
                                                                                                                                                                                                                                                                                                                                                                                                           location;
behaviour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                     CENT NAT
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                                                                                                                                                                                                                                                                98US-0108193P
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                                                                                                                                                                                                                                                                                                                                                                                                           gene marker;
; pedigree; s:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                     RECH SCI.
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Pred. No. 1.2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or nucleic acid encoding
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                                                                                                                                  genome, useful
behavioral traits
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                                            dog (Canine
selected
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The primers
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RESULT 264
AAF33001
ID AAF33
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                                                       Query Match
Best Local &
Matches 12
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Best Local 9
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                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                 Novel compound for diagnosing, preventing and treating immune disorders, comprising an oligonucleotide that specifically hybridizes with a nucleic acid sequence encoding B7 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (i.e. Type I and Type II) of markers. The map and the dog genome markers (or complementary sequences) are especially useful to identify genes responsible for phenotypic and behavioural traits in dogs, to identify morbid genes, to analyse diseases and identify implicated genes in such diseases and their alleles, and to study dog pedigrees. They may also be useful for isolating corresponding human gene sequences e.g. genes involved in genetic diseases
                                                                                                                                              dermatitis,
                                                                                                                                                        multiple sclerosis, psoriasis, systemic lupus erythematosus,
                                                                                                                                                                   the treatment of inflammatory and autoimmune disorders, including asthma juvenile diabetes mellitus, myasthenia gravis, Graves' disease, rheumatoid arthritis, allograft rejection, inflammatory bowel disease,
                                                                                                                                                                                                             The present invention provides sequences of antisense oligonucleotides targeted at the murine and human B7-1 and B7-2 coding and mRNA sequences. The antisense sequences have phosphorothicate backbones and some nucleotides are 2'-methoxyethoxy residues. The sequences can be used in nucleotides are 2'-methoxyethoxy residues. The sequences can be used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200074687-A1
                                                                                                                                                                                                                                                                                         Example 14; Page 84; 162pp; English
                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-049991/06.
                                                                                                                                                                                                                                                                                                                                                                                                               Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAY-2000; 2000WO-US014471.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human B7-2 antisense oligonucleotide SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-MAR-2001
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                            247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mouse; B7-1; B7-2; antisense; PCR primer; inflammation;
mune disorder; phosphorothioate backbone; ss.
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                                                                     Similarity
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                         TCCTGGAGCCCC 258
TCCTGGAGCCCC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGAAGGGCTGCA 6
                                                       Conservative
                                                                                                              BP; 3 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                           rhinitis, allergies and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                            Vickers TA,
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                                                                                 2.0%;
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Pred. No.
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Pred. No.
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hes 0;
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SASSESSES C

Mammalian; forensic; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; telomere metabolism; mutant; acquired immunodeficiency syndrome.

disease;

Phosphorothicate (PS) cligomer IV used to inhibit telomerase activity.

(first entry)

RESULT 266 AAS15460

AAS15460

standard;

DNA;

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TCCTGGAGCCCC

17

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247 TCCTGGAGCCCC 258

Matches Query Match

Local

l Similarity 12; Conserv

Conservative

0

Score 12; DB; Pred. No. 1.2
0; Mismatches

DB 4; L, 1.2e+05;

Length 20

Indels

0

Gaps

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2.0%;

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RESULT 265
AAF32817
Sequence
                                     nucleotides are 2'-methoxyethoxy residues. The sequences can be used in the treatment of inflammatory and autoimmune disorders, including asthy juvenile diabetes mellitus, myasthenia gravis, Graves' disease, rheumatoid arthritis, allograft rejection, inflammatory bowel disease, multiple sclerosis, psoriasis, systemic lupus erythematosus, contact
                                                                                                                                                                                           Novel compound for diagnosing, preventing and treating immune disorders, comprising an oligonucleotide that specifically hybridizes with a nuclei acid sequence encoding B7 protein.
                                                                                                          The present invention provides sequences of antisense oligonucleotides targeted at the murine and human B7-1 and B7-2 coding and mRNA sequence. The antisense sequences have phosphorothicate backbones and some
                                                                                                                                                                  Example 1; Page 50;
                                                                                                                                                                                                                                                     WPI; 2001-049991/06.
                                                                                                                                                                                                                                                                                                                                     04-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                              25-MAY-2000; 2000WO-US014471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human B7-2 mRNA antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF32817;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             autoimmune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-MAR-2001
                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mouse;
                                                                                                                                                                                                                                                                                Ġ
20 BP; 2 A; 7 C; 7 G; 4 T; 0 U; 0 Other;
                           rhinitis, allergies and cancer
                                                                                                                                                                                                                                                                               Vickers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B7-1;
                                                                                                                                                                                                                                                                                                                                     99US-00326186
                                                                                                                                                                  162pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B7-2; antisense; PCR primer;
                                                                                                                                                                                                                                                                            TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            phosphorothioate backbone; ss
                                                                                                                                                                                                                                                                               Karras
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RESULT 267
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                            individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFIP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other
                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of
                         14-FEB-2002
                                                                                                                                                                                                              Sequence 20 BP; 5 A; 6 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                             diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence represents a phosphorothicate (PS) oligomer used to inhibit telomerase activity in the methods of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New peptide nucleic acids that hybridizes to the RNA component mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative disease, autoimmune disease,
PNA XIV inhibiting human and mammalian telomerase activity
                                                                  AAS15455 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Col 29; 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-638024/73.
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09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunosuppressive;
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                                                                                                                                              з19 АССАТСТТСАСС 330
                                                                                                                                                                    1 Similarity
12; Conserv
                                                                                                                          <u>ب</u>
                                                                                                                                                                                                                                      invention
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                                                                                                                                                                     Conservative
                      (first entry)
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97US-00838545
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disease, or
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a sequence of 6-25 mulleobases, that inhibit telomerase activity in ammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of telomerase and as inhibitors of telomerase activity, or to component (hTR) sequence, as well as in forensic identification of criminal component (hTR) sequence, as well as in forensic identification of criminal component (hTR) sequence, as well as in forensic identification of criminal component (hTR) sequence, as well as in forensic identification of criminal component (hTR) sequence, as well as in forensic identification of criminal component of unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, component inflammation, component inflammation, component inflammation, acquired diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, human immunodeficiency virus (HTV) infections, acquired component inflammation in telomerase component inflammation in telomerase component inflammatical pathologies, and other component inflammatical pathologies, and other component inflammation in telomerase component inflammation in telomerase component inflammation in telomerase of the properties of the inflammation in telomerase of the inflammatical pathologies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New peptide nucleic acids that hybridizes to the RNA component mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalian; peptide nucleic acid; probe; forensic; paternity testinuman telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to peptide nucleic acids (PNAs), come sequence of 6-25 nucleobases, that inhibit telomerase activity
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09-APR-1997;
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
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